

1 Title:

2 *The ONE Study: Evaluation of Regulatory Cell Therapy in Kidney*  
3 *Transplantation Using a Harmonized Trial Design*

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86 **Abstract**

87 *Background*

88 Using cell-based medicinal products (CBMPs) represents a state-of-the-art approach to reducing  
89 general immunosuppression in organ transplantation. Accordingly, *The ONE Study* Consortium tested  
90 multiple regulatory cell products in kidney transplant (KTx) trials. Here, we report primary analysis  
91 results for overall safety of regulatory CBMPs when combined with reduced immunosuppressive  
92 treatment in this first *ONE Study* publication.

93 *Methods*

94 Seven investigator-led single-armed trials were conducted internationally in living-donor KTx  
95 recipients (60 week follow-up). One single-arm trial, the Reference Group Trial (RGT, n=66),  
96 represents a “standard-of-care” group given basiliximab, tapered steroids, mycophenolate mofetil  
97 (MMF) and tacrolimus. Data from six non-randomized phase I/IIa cell therapy group (CTG) trials were  
98 pooled and analyzed, where patients (n=38) received one of six CBMPs containing regulatory T cells,  
99 dendritic cells or macrophages; patient selection and immunosuppression mirrored the RGT, except  
100 basiliximab induction was substituted with CBMPs and MMF tapering was allowed. The primary  
101 endpoint was biopsy-confirmed acute rejection (BCAR); adverse event (AE) coding and immune  
102 monitoring was centralized.

103 *Findings*

104 Standard-of-care immunosuppression in the RGT recipients resulted in a 12·1% BCAR rate (expected  
105 range: 3·2-18·0%). The 6 CBMPs for the parallel CTG trials were administered to a combined total of  
106 38 patients, with an overall BCAR rate of 15·8%. 15 CBMP-treated patients (39·5%) were successfully  
107 weaned from MMF and maintained on tacrolimus monotherapy. Combined AE data and BCAR  
108 episodes from all six CTG trials revealed no safety concerns versus the RGT. Remarkably, fewer  
109 episodes of infections were registered in CTG trials versus the RGT. CTG, versus RGT, patients showed  
110 no loss of TSDR demethylation, no increase of CD8<sup>+</sup> T<sub>EMRA</sub> cells, and a healthy control-like restoration  
111 of immune cell composition (e.g. marginal zone-like B cells).

112 *Interpretation*

113 Regulatory cell therapy is achievable and safe in living-donor KTx recipients, and produces fewer  
114 infectious complications, but comparable rejection rates in the first year.

115

116 *Funding*

117 *The ONE Study* was funded by the 7<sup>th</sup> EU Framework Programme.

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## Introduction

121  
122 Combinations of general immunosuppressive drugs have enabled the widespread application of life-  
123 saving organ transplantation today; however, transplant survival is limited and has plateaued over  
124 the last decade,<sup>1</sup> leaving the dilemma of needing to replace damaged transplanted organs in a world  
125 where not enough organs are available, while the morbidity and economic costs associated with life-  
126 long general immunosuppression accrue. To address this problem, the organ transplantation  
127 community is well-aware that new strategies are urgently needed to decrease our dependency on  
128 immunosuppressive drugs to prevent allograft rejection.<sup>2</sup> Indeed, international networks have been  
129 established with this explicit purpose in mind, notably including a series of European Union-funded  
130 programs and, in North America, the *Immune Tolerance Network*. Research from these expert  
131 networks, and from numerous research laboratories across the globe, consistently call for novel  
132 therapies that will reduce our reliance on “full” immunosuppression to prevent organ rejection. At  
133 least two general strategies have been considered, including a deletional approach based on  
134 establishment of donor bone marrow chimerism to reduce donor-reactive immune cells, and an  
135 immune regulation-based approach that takes advantage of regulatory cells or pathways that control  
136 immunity and restrain immune responses to autologous antigens.<sup>3</sup> Although protocols to create  
137 chimerism in organ transplant recipients have been trialed for more than a decade, finding  
138 conditioning regimens with acceptable toxicity and avoiding the problem of graft-versus-host disease  
139 has been a persistent obstacle. Regarding the second strategy of building immune regulation, a  
140 therapeutic means to augment these cellular networks has only recently come of age for clinical  
141 testing.

142 Regulatory cell therapy has emerged as one attractive therapeutic approach to establish immune  
143 regulation aimed at protecting organ allografts.<sup>4-6</sup> The overall principle of this approach is to expand  
144 specific regulatory immune cell populations *ex vivo* in the form of cell-based medicinal products  
145 (CBMPs) that can then be infused into transplant recipients. Towards this aim, a European Union-  
146 funded consortium called *The ONE Study* was initiated with the aim of developing a range of CBMPs

147 and to test those cell products in early-phase clinical trials. The six CBMPs developed and tested in six  
148 parallel cell therapy group (CTG) trials ( $\leq 12$  patients each) in *The ONE Study* included two polyclonal T  
149 regulatory (pTreg), two donor-antigen reactive Treg (darTreg), one tolerogenic DC (ATDC) and one  
150 regulatory macrophage (Mreg) cell products. Central to the concept of *The ONE Study* was that all  
151 CBMPs be tested using the equivalent patient population of living-donor kidney transplant (KTx)  
152 recipients that receive the identical background immunosuppressive treatment, placing testing of the  
153 six CBMPs on a directly comparable basis. Also fundamental to this study was that a larger Reference  
154 Group Trial (RGT) be conducted on an equivalent patient population using standard-of-care  
155 immunosuppression. While the RGT is not strictly a true control group due to inclusion of basiliximab  
156 in place of cell therapy, it serves two purposes. First, since we have applied our CBMPs under similar,  
157 but reduced, immunosuppression, the RGT provides a recognized standard-of-care benchmark to  
158 assess whether currently expected outcomes are generally attainable with regulatory cell therapy  
159 with less immunosuppression. Second, with a standard-of-care RGT, performance of centralized  
160 immune monitoring allows for reliable detection of potential immunological changes caused by cell  
161 therapy. Here, we present the special design, clinical data, safety results and immune monitoring  
162 data of the *ONE Study* RGT and combined CTG group of trials, which is intended as a foundation for  
163 further regulatory cell therapy trials in organ transplantation.

164

165

## Methods

### 166 Study design and participants

167 The *ONE Study* aimed to explore the safety and immunological effects of regulatory cell-based  
168 therapy as an adjunct immunosuppressive treatment in living-donor kidney transplant recipients  
169 through a series of clinical trials sharing the same general design. Therefore, we created a multi-trial  
170 design strategy to facilitate: 1) comparison of different cell therapy trials versus standard-of-care  
171 treatment, and 2) comparison of cell therapy trials to each other. In total, seven trials were  
172 performed, the first being the single-arm multi-center RGT conducted at all clinical sites that were  
173 planning to perform an individual cell therapy trial. The RGT formed the basis for the other six  
174 individual trials testing CBMPs (the CTG trials). Chronologically, enrollment for the RGT was  
175 completed before any of the CTG trials commenced; the RGT was initiated while regulatory approvals  
176 for the CTG trials and cell manufacturing procedures were being obtained.

177

178 *CBMPs*. In the course of *The ONE Study* project, six regulatory cell products were approved for  
179 manufacture and therapeutic testing in the CTG trials by the national competent authority in each  
180 participating country. Two of the six cell products consisted of polyclonal natural T regulatory cells  
181 approved respectively in the United Kingdom (“pTreg-1”)<sup>7</sup> and Berlin (“pTreg-2”)<sup>8</sup>. The third and  
182 fourth cell products consisted of Treg, but were generated in the presence of donor antigen during  
183 manufacturing; one product was exposed under conditions of costimulatory blockade in the  
184 presence of donor peripheral blood mononuclear cells (PBMCs) in Boston<sup>9</sup> (referred to as  
185 costimulatory blockade “darTreg-CSB”) and the other product was developed in San Francisco where  
186 Tregs sorted from PBMCs were stimulated with donor B cells that had been activated with K562 cells  
187 expressing human CD40L (referred to as donor alloantigen-reactive “darTreg-sBC”)<sup>10</sup>. The fifth and  
188 sixth cell products were derived from peripheral blood monocytes, where monocytes were  
189 stimulated in Nantes with GM-CSF to produce autologous tolerogenic dendritic cells (“ATDC”),<sup>11</sup> or in  
190 Regensburg with M-CSF and IFN- $\gamma$  to produce regulatory macrophages (“Mreg-UKR”)<sup>12</sup>. All six

191 regulatory cell products were derived from recipient leucocytes (blood or leucopheresates), with the  
192 exception that Mreg-UKR were donor-derived. Table S1 provides an overview of the overall  
193 characteristics of the CBMPs, including a reference to cell production methods.

194

195 *Patient selection for trials.* Living-donor KTx recipients were selected for inclusion into all seven trials.  
196 Living donors were chosen for these trials to allow for maximal planning logistics with regard to  
197 obtaining informed consent, having a medically stable recipient population, coordinating regulatory  
198 cell manufacturing from donor or recipient cells (in the CTG trials) and obtaining pre-transplant  
199 immune monitoring samples. The core inclusion and exclusion criteria that were common to all trials  
200 for both the donors and recipients are listed in Table S2. The main exclusion criteria were patients  
201 transplanted previously, high risk recipients (PRA >40%) and HLA identical donor-recipient  
202 mismatches (0-0-0 mismatches); all patients needed to be  $\geq 18$  years old.

203

204 *RGT treatment protocol.* The ONE Study group of clinicians developed the RGT immunosuppression  
205 design based on their own local standard-of-care protocols, which included some features of the  
206 ELITE-Symphony study<sup>13</sup>, for the selected non-high risk KTx patient population. The study protocol  
207 (clinicaltrials.gov: NCT01656135) consisted of: basiliximab administration  $\leq 2$  hours before transplant  
208 surgery and on day 4 after surgery (20mg i.v.); prednisolone starting on day 0 (day of KTx) and  
209 gradually tapered away by week 15; mycophenolate mofetil (MMF) at 2 g/day from day -1 to day  
210 +14 and 1.5 g/day thereafter; and tacrolimus starting on day -4 at 3-12 ng/ml and gradually reduced  
211 over 9 months to 3-6 ng/ml. A diagram showing the exact dosing scheme can be found in Fig. S1.  
212 Patient follow-up was continued for 60 weeks. The target recruitment figure for the RGT was 60  
213 patients.

214

215 *CTG treatment protocol.* The clinical protocol for the 6 CTG trials closely mirrors the regimen for the  
216 RGT (Fig. S1). All cell products were delivered once intravenously between day -7 and day +10



217 relative to the day of KTx; within this timeframe, monocyte-derived cell products were administered  
218 before KTx and T cell-derived products were given after KTx. The exact cell numbers infused will be  
219 provided in the individual CTG trial descriptions to be reported elsewhere, but ranged from 0.5 to 10  
220  $\times 10^6$  cells/Kg BW for all cell products except darTreg-CSB, where a range between  $2 \times 10^3$  -  $2 \times 10^6$   
221 cells/Kg BW was targeted. Pharmacological immunosuppression and dosing were the same as with  
222 the RGT, except that basiliximab induction therapy was omitted, and at 9 months post-KTx an option  
223 was included to completely taper away MMF by one year post-KTx; with MMF cessation, tacrolimus  
224 continued as a monotherapy. Tapering of MMF was not allowed if an immediately prior KTx biopsy  
225 showed signs of subclinical rejection or there was evidence of declining renal function. Patient  
226 follow-up continued for approximately 60 weeks, after which time immunosuppressive treatment  
227 was decided by the local transplant physician. The number of cell therapy-treated patients did not  
228 exceed 12 in any individual CTG trial. All CTG trials are registered on ClinicalTrials.gov (NCT02252055,  
229 NCT02085629, NCT02244801, NCT02371434, NCT02129881 and NCT02091232).

230

231 *Sites performing trials.* The multicenter RGT was performed at eight international locations, including  
232 the University Hospital Regensburg (Regensburg, Germany), Charité (Berlin, Germany), Centre  
233 Hospitalier Universitaire Nantes (Nantes, France), Ospedale San Raffaele (Milan, Italy), Oxford  
234 University Hospitals NHS Foundation Trust (Oxford, UK), Guy's Hospital (London, UK), Massachusetts  
235 General Hospital (Boston, MA) and UCSF Medical Center (San Francisco, CA) (Fig. 1). After completing  
236 enrollment for the RGT, seven centers conducted a separate CTG trial with one of six regulatory cell  
237 products (see above). Unlike the five centers that recruited patients into their respective single-  
238 center CTG trials, the Oxford and London sites joined forces to recruit patients into one CTG trial  
239 (pTreg-1). Notably, the Milan site participated only in the RGT, since their cell product was not  
240 approved for clinical trial testing during *The ONE Study*.

241

242 *Endpoints.* Biopsy-confirmed acute rejection (BCAR) was the primary endpoint. Histopathological  
243 grading of KTx biopsies was performed by a central pathologist (Prof. Ian Roberts, Oxford University)  
244 for all trials within *The ONE Study*, with the standard assessment performed according to the Banff  
245 criteria.<sup>14</sup> Notably, a case of borderline histological change in a for-cause biopsy with clinical evidence  
246 of acute rejection was considered a BCAR. However, histological changes consistent with acute  
247 rejection that were not accompanied by clinical evidence of rejection were not recorded as a BCAR,  
248 but were logged as a secondary endpoint. Estimated glomerular filtration rate (eGFR: MDRD method)  
249 was recorded as a secondary endpoint.

250

251 For the RGT, we estimated a BCAR rate of approximately 10% after 60 weeks under standard  
252 immunosuppressive therapy in the select KTx patient population. With this assumption, a two-sided  
253 95% confidence interval for a single proportion of 0.106 predicts a rejection rate ranging from 3.2-  
254 18.0% with a sample size of 66 patients; a BCAR rate falling outside this interval would suggest that  
255 the rejection rate is atypical.

256

257 *Clinical data collection and monitoring.* Clinical data from all trials were entered into a web-based  
258 data capture platform consisting of electronic case report forms (eCRF) custom-made for *The ONE*  
259 *Study* (Koehler eClinical, Freiburg, Germany). A core set of clinical data were collected from all trials  
260 to ensure that these parameters could be directly compared. Selected data items for evaluation of  
261 the study endpoints were verified for accuracy against source documents during on-site monitoring  
262 visits performed by qualified CRAs. Additionally, data were reviewed, queried and cleaned remotely  
263 by a central team of data managers using both automatic and manual data validation checks. All  
264 adverse events (AEs) and serious adverse events (SAEs) were coded centrally using version 20.1 of  
265 the Medical Dictionary for Regulatory Activities (MedDRA) and quality-controlled to ensure  
266 consistency of coding across all trials and study sites. To compare safety events reported from  
267 cohorts of different sizes, (S)AE frequencies were normalized using a cohort-specific "Patient Study

268 Years” (PSY) denominator. PSY is the cumulative amount of time spent by trial participants in study  
269 follow-up and was calculated and applied for RGT and CTG separately. A safety advisory board (SAB)  
270 received SAE reports for all CTG trials as they occurred and reviewed all safety data twice per year.  
271 To be sure of open communication within *The ONE Study* trial series, safety alerts or conclusions  
272 from the SAB were shared with all centers performing CTG trials.

273

274 *Immune monitoring.* We used a mixed model of locally and centrally performed assays to compare  
275 pre- and post-transplant immune status of RGT and CTG trial patients.<sup>15</sup> The following analyses were  
276 performed as provided in supplementary materials: immune cell composition by whole blood flow  
277 cytometry, TSDR demethylation **gene expression (see Supplementary Methods)** and anti-donor as  
278 well as anti-CMV IFN $\gamma$  EliSpot. **To reveal differences in peripheral blood immune cell composition**  
279 **between patients with end-stage renal disease (RGT and CTG before transplantation) and healthy**  
280 **individuals, we performed comparative analyses with age-and gender-matched healthy controls from**  
281 **our recently generated cohort data set.**<sup>16</sup>

282

283 *Statistical analyses.* **A statistical analysis plan defined the conventions and analyses, and emphasized**  
284 **the exploratory nature of the *ONE Study*, accordingly the proposed statistical examination of clinical**  
285 **data was descriptive. The reported comparative analyses of changes in immune cell composition and**  
286 **functionality between RGT and CTG patients were done as *post-hoc* analyses.**

287

288 For clinical data, results for baseline characteristics, safety and transplant function or rejection  
289 endpoints were summarized descriptively. **No formal testing was performed. In addition to crude**  
290 **rejection rates, time to first BCAR was analyzed using Kaplan-Meier methods. The primary BCAR**  
291 **endpoint is reported descriptively for the intention-to-treat population (RGT, n=66; CTG, n=38); the**  
292 **time-to-event Kaplan-Meier BCAR analysis is presented for both the intention-to-treat (66/38,**  
293 **respectively) and per-protocol (47/32, respectively) populations. All other variables (DSA, eGFR,**

294 tacrolimus levels) are summarized for the number of patients who were tested at the relevant study  
295 time points. Incidence rates of adverse events normalized per 100 patient study years were  
296 calculated and based on the intention-to-treat population.

297

298 Differences in immune monitoring results between RGT patients prior to transplantation and healthy  
299 controls were analyzed applying Kruskal Wallis tests followed by Dunn-Bonferroni tests. Changes  
300 between pre-transplant and post-transplant time points of the same patient were analyzed applying  
301 Wilcoxon matched-pairs signed rank test. To reveal differences in immune cell composition or TSDR  
302 changes after transplantation between RGT and CTG patients, we employed a Kruskal Wallis and a  
303 post-hoc Dunn's multiple comparison test. P values <0.05 were considered as significant.

304

305 *Role of the funding source.* The funders had no role in data collection, analysis, interpretation or  
306 writing of the manuscript. EKG, as the ONE Study Consortium FP7 project coordinator, had access to  
307 all the data in the study; BS also had access to the full data set. As a group, members of this FP7  
308 consortium discussed the publication plans, and therefore were involved in the decision to submit  
309 the manuscript; EKG and BS had the final responsibility in this decision.

## Results

310

### 311 Results from clinical trials

312 *RGT and CTG trials conduct.* Recruitment to the RGT began in December 2012, with the last patient-  
313 last visit in December 2015. Fig. 1 shows that 70 patients were enrolled in the RGT in total (red arrow  
314 bars), with 66 receiving a KTx. Of the four pre-KTx withdrawals, two had their transplant postponed,  
315 one patient needed treatment for DSA that did not allow further inclusion into the study protocol,  
316 and one patient withdrew consent. 61 RGT patients completed the study: of the five who were non-  
317 completers, one patient withdrew consent (at 8 days), one patient was lost to follow-up (at 33  
318 weeks), one patient had a major vascular complication and graft loss (at 8 days), one patient received  
319 ATG instead of basiliximab induction therapy (discovered on day 11), and one patient violated the  
320 eligibility criteria (noted at 24 weeks). None of these five patients registered a primary endpoint. In  
321 the RGT, median follow-up time was 60.1 weeks (IQR 1.3 weeks). Fig. 1 also summarizes patient  
322 recruitment into the six individual CTG trials (non-red arrow bars), where a total of 60 patients were  
323 recruited into the various trials, with the first patient-first visit conducted in May 2014 and the last  
324 patient-last visit done in November 2018. Of the 60 enrolled patients, 38 received a KTx and the  
325 designated cell therapy. All of these patients completed the 60 week follow-up planned in the *ONE*  
326 *Study*. The 22 patients withdrawn were due to one of the following: cell manufacturing failures (14),  
327 early development of acute rejection before the planned cell infusion (5), discovery of ineligibility  
328 criteria after enrollment (2) or requirement for a second abdominal surgery shortly after KTx (1). Cell  
329 manufacturing failures were because of failure to meet release criteria (9), cancellation (2),  
330 microbiology testing positive (2) and leucapheresis side effects (1); no trial was stopped due to lack  
331 of manufacturing feasibility. In the CTG, median follow-up time was 60.0 weeks (IQR 0.6 weeks). A  
332 summary of the recipient and donor demographic data for the RGT and CTG trials is provided (Tables  
333 S3 and S4). Data on recipient and donor age, gender, ethnicity, renal replacement therapy,  
334 relationship of donor and recipient, and underlying diagnosis show that the RGT and combined CTG  
335 trials were well-balanced. Notably, both the RGT and combined CTG trials have a nearly identical

336 over-representation of male recipients; since gender-related effects are known in transplantation,  
337 this should be taken into consideration when interpreting the results.

338

339 A set of per protocol criteria were defined based mostly on overall adherence to the planned  
340 immunosuppression regime in both the RGT and CTG trials (criteria listed in Table S5). In the RGT, 47  
341 of 66 KTx patients (71.2%) received treatment that closely followed the clinical protocol, whereas 32  
342 of the 38 patients (84.2%) in the CTG trials were treated with close adherence to the protocol.

343 Reasons for non-adherence varied widely among the trials, but were mostly related to adjustments  
344 or switching of immunosuppression that the treating physician deemed necessary. Furthermore, *ONE*  
345 *Study* physicians performing the CTG trials tapered immunosuppression to tacrolimus monotherapy  
346 (optional) in 17 of 38 (44.7%) patients. The immunosuppression was successfully tapered in all but  
347 two cases, where triple therapy was later reinstated due to a BCAR and detection of recurrent IgA  
348 nephropathy, respectively.

349

#### 350 **Outcomes (BCAR rate, GFR, DSA, tacrolimus levels)**

351 BCAR rate in the RGT was 12.1% (8/66), which is within the expected range of 3.2-18.0%. BCAR  
352 occurred in 15.8% (6/38) of the patients receiving cell therapy within the combined CTG trials, which  
353 was within the expected range calculated for the RGT. The Kaplan-Meier curves in Fig. 2A highlight  
354 the early incidence of BCARs in all trials. The severity of the first BCAR by Banff scoring was  
355 distributed similarly between the RGT and the group of CTG trials (Fig. 2B); one patient in the RGT  
356 experienced a second BCAR episode, but other BCARs in all trials were single episodes and were  
357 successfully treated. Only one of eight first BCAR episodes in the RGT occurred after two weeks post-  
358 KTx; similarly, 4 of 6 episodes of BCAR in the CTG group trials occurred before three weeks post-KTx.  
359 Specific BCAR data from individual sites will be published separately for each CTG trial. In addition,  
360 we also performed a Kaplan-Meier analysis for the “per protocol” patients in the RGT and group of  
361 CTG trials (Fig. 2C); the rate and timing of the BCAR episodes were essentially the same.

362

363 A set of tests was performed at study end (60 weeks) to further assess outcomes in the trials,  
364 including DSA detection, eGFR and tacrolimus blood levels. At study end, DSA testing revealed that  
365 13.7% (7/51 tested) of RGT recipients had a DSA, with 15.2% (5/33 tested) showing DSA in the  
366 combined CTG trials; of the CTG patients tapered to monotherapy, 13.3% (2/15 tested) had a new  
367 DSA. Regarding kidney function (Fig. S2), eGFR measurements in the RGT and CTG trials showed an  
368 almost identical increase over the study period (20.4% and 20.8%, respectively) when comparing  
369 median eGFR at 60 weeks post-KTx to median eGFR at one week post-KTx. As a reflection of  
370 immunosuppressive load at study end, tacrolimus trough levels were found to be similar in the RGT  
371 and combined CTG trials, at  $6.1 \pm 2.1$  (mean $\pm$ SD; n=44 tested) and  $6.6 \pm 1.6$  ng/ml (mean $\pm$ SD; n=32  
372 tested), respectively. Furthermore, immunosuppressive burden with tacrolimus (trough level: Fig.  
373 S3A, B) and MMF (dose: Fig. S3C, D) was similar or even tended to be lower throughout the study  
374 period in the CTG versus RGT patients. Together, these data should be considered with the  
375 understanding that 15 patients (39.5%) in the CTG trials were on tacrolimus monotherapy at study  
376 end, whereas 98.4% (60/61) of patients in the RGT continued on at least dual immunosuppression.

377

### 378 **Safety Data**

379 The normalized incidence rates of treatment-emergent SAEs/AEs in the RGT (n=66) and CTG trials  
380 (n=38) were 91.2/1614.6 and 70.7/1452.0 events per 100 PSY, respectively, indicating no increase in  
381 adverse events with cell therapy (Table S6). In the CTG trials, there was special attention given to  
382 identifying SAEs/AEs related to cell therapy infusion. Overall, there were 12 AEs reported with a  
383 possible relationship to the cell infusion, only one of which was a serious incident (an SAE; increased  
384 creatinine) (Table S7). All potentially related adverse events only occurred once, so no specific  
385 pattern was exposed in the 38 patients treated with CBMPs. No deaths were reported in any of the  
386 trials.

387

388 A descriptive analysis of normalized data comparing MedDRA-coded SAEs in the RGT versus the  
389 combined data from the CTG trials revealed that most serious medical problems were similar in  
390 frequency (Fig. 3A). However, there was one substantial difference that emerged which is worth  
391 considering in detail. The incidence rate of SAEs in the RGT related to infections and infestations was  
392 nearly six-fold higher compared to the combined CTG trials. After examining all infection-related  
393 adverse events (AEs) recorded in the trials, this pattern of decreased infections in the CTG trials was  
394 consistently observed across the CTG trials (Fig. 3B) and was evident during the entire post-KTx  
395 observation period (Fig. 3C). Also interestingly, we found that the main difference was with regard to  
396 a reduced number of viral infections in the CTG trials (Fig. 3D); notably, there was also an appreciable  
397 difference in the number of infections recorded without specifying the pathogen, but numbers of  
398 bacterial and fungal infections were essentially the same. Breaking the data down even further  
399 regarding AEs, the main decreases in viral infections in the CTG trials were with regard to CMV,  
400 herpes (including herpes simplex, herpes-zoster, oral herpes, nasal herpes and Varicella-zoster) and  
401 polyoma virus (Fig. 3E). The decreased rate of viral infection in the CTG was not due to more  
402 preventive measures, since 65.2% (43/66) RGT and 52.6% (20/38) CTG patients received anti-viral  
403 prophylaxis in the first three months after KTx; also, notably, the percentage of CMV<sup>+</sup> to CMV<sup>-</sup> donor  
404 to recipient transplants was 18.2% and 21.1% in the RGT and CTG trials, respectively. Therefore,  
405 patients receiving cell therapy in general developed fewer viral infections compared to patients  
406 receiving standard-of-care treatment.

407

#### 408 **Immune monitoring results**

409 Identical standardized immune monitoring testing of peripheral blood cells was performed in all  
410 patients of the seven trials. In general, principal component analyses show that RGT patients prior to  
411 KTx have major alterations in absolute and relative blood immune cell population composition  
412 compared to age- and gender-matched healthy controls (Fig. 4A). Populations contributing most to  
413 those alterations were granulocytes, CD16<sup>+</sup> mDCs and CD14<sup>high</sup>CD16<sup>+</sup> intermediate monocytes, which



414 were increased in RGT patient samples, but also plasmacytoid DCs (pDCs), marginal zone-like B cells  
415 (MZB) and CD8<sup>+</sup>CD28<sup>+</sup> T cells which were higher in samples of healthy controls (Fig. 4B). Post-KTx  
416 longitudinal analysis revealed only moderate or absent normalization of CD16-expressing monocytes  
417 and MZB, respectively (Fig. 4C). Furthermore, whereas composition of conventional CD4<sup>+</sup> T cells  
418 subsets remained normal and comparable to healthy controls, CD8<sup>+</sup> T cells subset composition showed  
419 major alterations over the post-KTx course. Although naïve T cells increased early after transplantation,  
420 we observed a skewing towards terminal differentiation of CD8<sup>+</sup> T cells in the long-term (Fig. 4C).

421

422 Examining immunophenotyping results from the RGT and combined CTG trials, we did not observe  
423 significant differences in numbers or proportions of CD4<sup>+</sup>CD25<sup>high</sup>CD127<sup>low</sup> Tregs between the groups  
424 at 15 months post-KTx (Fig. 5A). A significant reduction in TSDR demethylation occurred in RGT  
425 patients, but not in CTG trial patients. Furthermore, only RTG patients showed a significant increase in  
426 CD8<sup>+</sup> T<sub>EMRA</sub> cells and CD8<sup>+</sup>CD57<sup>+</sup> chronically-activated T cells (Fig. 5B), whereas in samples from CTG  
427 patients we observed more CD8<sup>+</sup>CD28<sup>+</sup> T cells. Both patient groups showed a reduction of donor-  
428 specific IFN $\gamma$  producing memory T cells after KTx (Fig. S4A). However, RGT patients in contrast to CTG  
429 patients showed higher anti-CMV T cell responses (Fig. S4B), which correlated with absolute CD8<sup>+</sup> T<sub>EMRA</sub>  
430 numbers (Fig. S4C). This increase is well known in KTx patients and is likely related to inflammation  
431 triggered subclinical reactivation of CMV, which we also only observed in RGT but not CTG patients  
432 (Fig. 3E). Although both patient groups had more pDCs 15 months post-KTx, we only observed a  
433 normalization of MZB numbers and a significant reduction of CD14<sup>high</sup>CD16<sup>+</sup> monocytes in CTG patients  
434 (Fig. 5C). In addition, CTG patients showed increased mRNA expression of genes described to be high  
435 in immunosuppression-free operationally tolerant kidney transplant patients (e.g. Ms4A1) and co-  
436 inhibitory molecules (CD200), but reduced expression of rejection-associated genes (HMMR, Fig. S4D).  
437 Together, these data suggest that regulatory cell therapy within our trials CTG patients show a more  
438 healthy control-like restoration of immune cell composition.

439

440

## Discussion

441 The *ONE Study* consortium has taken the unique approach of performing side-by-side trialing of  
442 different T cell, DC and macrophage regulatory cell products in low to medium risk KTx recipients. In  
443 this coordinated group of six international early phase clinical trials (the CTG trials), we show that  
444 CBMP application in this patient population is feasible for multiple regulatory cell types, and their  
445 categorical application near the time of KTx reveals no apparent safety concerns, including allograft  
446 rejection rate. Furthermore, 15 of the 38 patients treated with CBMPs were successfully weaned to  
447 tacrolimus monotherapy during the 60 week observation period. The conduct of a parallel reference  
448 trial (the RGT) by the same clinical sites collecting matching clinical information and immune  
449 monitoring data provided a standard-of-care benchmark to confidently assess critical safety and  
450 immunological parameters, and also to evaluate whether reduction of immunosuppression through  
451 CBMP application could have potential benefits to patients. Remarkably, in this regard, the rate of  
452 viral infections was considerably lower in patients treated with regulatory cell products compared to  
453 standard-of-care treatment, particularly with regard to viral infections. Furthermore, centralized  
454 immune monitoring of peripheral blood leucocyte populations suggests a return of CBMP-treated  
455 (CTG), but not conventionally-treated (RGT), recipients towards a state of immune homeostasis.  
456 Therefore, results from the *ONE Study* establish a fundamental basis for further testing of regulatory  
457 cell CBMP therapy in organ transplantation, and provide initial evidence that reducing general  
458 immunosuppressive burden through cell therapy could potentially decrease serious side effects in  
459 KTx recipients.

460

461 This initial *ONE Study* report focusses only on the CTG trials as a combined group, and not on results  
462 from the individual CTG trials. While each of the six individual CTG trials followed the same clinical  
463 treatment protocol with regard to background immunosuppression, thus allowing for a  
464 comprehensive analysis of the CTG trials as a whole group, there are important details from each of  
465 those trials that deserve in-depth reporting and explanations in additional follow-up publications.

466 Indeed, forthcoming details from the individual cases will provide insight into interesting feasibility,  
467 safety aspects and effects of each specific cell therapy product, permitting examination of issues such  
468 as cell production methods, CBMP characterization, cell dosing, infusion scheduling, clinical  
469 outcomes and immunological features from KTx biopsy specimens, as well as a comprehensive set of  
470 central immune monitoring results. Nonetheless, the current analysis of results from the combined  
471 CTG trials provides a uniquely broad evaluation of safety and outlook perspective for cell therapy in  
472 organ transplantation, and shows that cell therapy was feasible in terms of logistics and cell  
473 manufacturing in the majority (38/52: 73%) of patients ready to receive the therapy.

474

475 One of the main motivations for seeking new therapies in organ transplantation is to reduce the  
476 need for general immunosuppressive drugs, which have substantial toxicities and incrementally  
477 expose recipients to dangers inherent from a suppressed immune system, most commonly  
478 infections. A recent set of guidelines and comprehensive review by Fishman<sup>17</sup> highlights the extent of  
479 the infection problem, and its direct relationship to immunosuppressive load. Results from the *ONE*  
480 *Study* CTG trials indicate that lowering immunosuppression does appear to decrease the risk for viral  
481 infections. This was also supported by the immune monitoring results, as only RGT patients showed a  
482 tendency towards increased proportions of CMV-specific memory T cells correlating with signs of  
483 chronic CD8<sup>+</sup> T cell activation at the end of the observation period, as previously described.<sup>18-20</sup> What  
484 remains unknown at this point is whether decreased infections were simply due to less  
485 immunosuppression in the CTG trials, or were related in some way to the cell therapy action itself;  
486 neither possibility can be ruled out. However, by keeping the overall inflammation low, regulatory  
487 cell therapy may prevent reactivation of persistent viruses such as CMV and other herpes viruses. It  
488 should be noted that immunosuppressive burden was lower early-on post-KTx (no basiliximab  
489 induction) and in some patients after nine months post-KTx (MMF tapering), but that the infection  
490 rates were consistently less across the spectrum of CTG trials during the entire observation period  
491 (Fig. 3C). While reduction of MMF treatment is within the prophylactic guidelines for patients at risk

492 for developing viral infection,<sup>17</sup> the gap in reported infections did not show evidence of widening  
493 between the RGT and CTG trials after nine months, leaving this issue an open question. Nonetheless,  
494 our data encourage the performance of prospective randomized clinical trials to confirm an  
495 infectious disease benefit from regulatory cell therapy protocols.

496

497 Our immune monitoring results showed that patients with end-stage renal failure exhibit major  
498 alterations in their peripheral immune cell composition compared to age- and gender-matched  
499 healthy controls, most likely reflecting their increased inflammatory state.<sup>21-23</sup> Standard  
500 immunosuppressive therapy in RGT patients did not reverse these alterations, but rather led to  
501 further immune cell imbalance as evidenced by a significant reduction in markers for stable Tregs.<sup>24</sup>  
502 Importantly, regulatory cell therapy mitigated this Treg reduction and correlated with a healthy  
503 control-like restoration of immune cell composition. In particular, MZB numbers, also discussed to  
504 have anti-inflammatory or regulatory function,<sup>25,26</sup> were increased in CTG patients at the end of the  
505 observation period. Thus, although both RGT and CTG trial patients had a reduction in donor-specific  
506 IFN $\gamma$ -producing memory T cells, only the cell therapy-treated patients tended to experience a re-  
507 establishment of immune cell homeostasis, which is a major goal in organ transplantation.

508 **Importantly, these immune-related differences were independent of potential confounding factors**  
509 **such as donor relationships.** Whether this effect is related to cell therapy itself, or is due to reduced  
510 immunosuppressive load in the CTG trials, will need to be investigated further in future trials.

511

512 To date, there are few published reports on the use of regulatory cell therapy in human organ  
513 transplantation, some of which were pilot trials conducted previously by *ONE Study* investigators  
514 [recently reviewed by Romano 2019]. Hutchinson and colleagues have tested different preparations  
515 of regulatory macrophages in KTx recipients,<sup>27-29</sup> which provided critical lessons for designing the  
516 *ONE Study* CTG trials. Additionally, polyclonal Tregs have been administered by the UCSF group to  
517 three KTx recipients with biopsy-proven subclinical inflammation six months after transplantation,

518 showing that cell therapy is feasible in this circumstance;<sup>30</sup> late administration of expanded  
519 polyclonal Tregs has also been reported by the Northwestern group in nine lymphodepleted KTx  
520 recipients.<sup>31</sup> In liver transplantation, Todo et al. have infused costimulatory blockade conditioned  
521 lymphocytes similar to those used by the MGH group in the *ONE Study*, and were able to achieve  
522 complete immunosuppression withdrawal in seven of the ten splenectomized and  
523 cyclophosphamide-conditioned recipients.<sup>32</sup> Unfortunately, these pilot studies are highly variable in  
524 design, and did not incorporate a parallel trial with a similar group of patients not receiving cells to  
525 better appraise whether cell therapy is safe or shows indications of discernable effects. Importantly,  
526 the *ONE Study* trials were developed with the fundamental viewpoint that a reference trial, and also  
527 comparison to healthy control data, is absolutely necessary to make practical conclusions about  
528 regulatory cell therapy testing. Therefore, to advance the cell therapy field in organ transplantation,  
529 we aimed to evaluate cell therapy against a recognized standard-of-care (RGT) treatment by infusing  
530 different CBMPs near the time of KTx as a replacement for conventional induction treatment  
531 (omitting basiliximab induction). Into this design we incorporated an option to wean MMF starting at  
532 nine months to further offer potential benefit to patients from general immunosuppression, and to  
533 stress-test this cell therapy protocol under rigorous clinical monitoring. With this overall study  
534 strategy, and by performing the RGT as a multicenter study together with the CTG trials as parallel  
535 individual trials at the same sites, the *ONE Study* consortium uniquely delivers meaningful and  
536 reliable information about regulatory cell therapy to the organ transplantation community. Based on  
537 the *ONE Study*, the UK group has already initiated a randomized trial called the *TWO Study* with their  
538 polyclonal Treg cell product (ISRCTN11038572), and other *ONE Study* partners (Massachusetts  
539 General Hospital: NCT03577431 and UCSF Medical Center: NCT02188719) are conducting trials in  
540 transplant recipients with cell products used in the *ONE Study*. Opening the way to these and other  
541 more advanced clinical trials was the unifying philosophy of the *ONE Study*.  
542

543 **Contributors**

544 BS, EKG, PNH, PR, AM, JAH, DSG, CAB, JFM, TB, RC, WP, NA, HJS, PJF, RH, MB, BJ, JBN, MPH-F, UK,  
545 SJK, JG, PJM, LB, LAT, RJL, AB, JAB, GL, KJW, MCC, AS, BB, GB, SMK, and HDV contributed to the study  
546 design. PNH, PR, JAH, DSG, CAB, JFM, TB, RC, WP, NA, HJS, PJF, RH, JBN, AS, BB, GB, SMK, NMO, and  
547 RÖ managed patient care. PR, AM, JAH, MB, AB, JAB, GL, KJW, MCC, QT, CS, ECG, LC-R, KC, ME, SK,  
548 and AS were involved in cell production. BS, EKG, PNH, PR, AM, JAH, MB, BJ, JBN, MPH-F, AB, MCC,  
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551 MS, RJ, CM, and SS performed data analysis. BJ, CM, SS, and KJ were study statisticians. BS, EKG, PNH,  
552 PR, UK, SJK, JG, PJM, LB, LAT, RJL, AB, JAB, GL, KJW, MCC, AS, BB, GB, SMK, HDV, AS, ISDR, MS, RJ,  
553 CM, SS, and KJ interpreted data. EKG and BS wrote the manuscript, which was reviewed by JAH, BJ,  
554 SS, and KJ, as well as the other authors. EKG was the *ONE Study* EU FP7 project coordinator.

555

556 **Declaration of interest**

557 BS, PR, AM, JAH, DSG, QT, ECG, MB, WJB, ISDR, MS, RJ, JFM, CB, BJ, LC-R, RC, IM, NMO, MPH-F, CM,  
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573 20150110761 A1 issued and is a founder and current CEO of Sonoma Biotherapeutics which works  
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580

581 **Data sharing**

582 We will follow the common controlled access principles outlined by the Medical Research Council  
583 Clinical Trials Unit (<https://www.ukri.org/funding/information-for-award-holders/data-policy/>).

584 According to those principles, we will acknowledge that data with long-term value be preserved, and  
585 usable for future research. We do, however, want to ensure that there are legal, ethical and  
586 commercial constraints maintained on the release of research data according to the following code.  
587 Research teams are entitled to receive appropriate recognition for their efforts in collecting and  
588 analyzing data and should be given at least a limited period of privileged to use and publish the data,  
589 before key trial data are open for use by other researchers. If such requests are made to access the  
590 data, resources need to be available in order to process the request and prepare the data in a timely  
591 manner, if possible. Because of these demands, there must be an important scientific objective  
592 behind each request. Especially in the case our internationally conducted *ONE Study*, any request  
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595

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606 **References:**

- 607 1. Wekerle T, Segev D, Lechler R, Oberbauer R. Strategies for long-term preservation of kidney  
608 graft function. *Lancet* 2017; **389**(10084): 2152-62.
- 609 2. Bamoulid J, Staeck O, Halleck F, et al. The need for minimization strategies: current problems  
610 of immunosuppression. *Transpl Int* 2015; **28**(8): 891-900.
- 611 3. Rickert CG, Markmann JF. Current state of organ transplant tolerance. *Curr Opin Organ*  
612 *Transplant* 2019; **24**(4): 441-50.
- 613 4. Safinia N, Grageda N, Scotta C, et al. Cell Therapy in Organ Transplantation: Our Experience  
614 on the Clinical Translation of Regulatory T Cells. *Front Immunol* 2018; **9**: 354.
- 615 5. Marin E, Cuturi MC, Moreau A. Tolerogenic Dendritic Cells in Solid Organ Transplantation:  
616 Where Do We Stand? *Front Immunol* 2018; **9**: 274.
- 617 6. Hutchinson JA, Geissler EK. Now or never? The case for cell-based immunosuppression in  
618 kidney transplantation. *Kidney Int* 2015; **87**(6): 1116-24.
- 619 7. Fraser H, Safinia N, Grageda N, et al. A Rapamycin-Based GMP-Compatible Process for the  
620 Isolation and Expansion of Regulatory T Cells for Clinical Trials. *Mol Ther Methods Clin Dev* 2018; **8**:  
621 198-209.
- 622 8. Landwehr-Kenzel S, Zobel A, Hoffmann H, et al. Ex vivo expanded natural regulatory T cells  
623 from patients with end-stage renal disease or kidney transplantation are useful for autologous cell  
624 therapy. *Kidney Int* 2018; **93**(6): 1452-64.
- 625 9. Guinan EC, Cole GA, Wylie WH, et al. Ex Vivo Costimulatory Blockade to Generate Regulatory  
626 T Cells From Patients Awaiting Kidney Transplantation. *Am J Transplant* 2016; **16**(7): 2187-95.
- 627 10. Putnam AL, Safinia N, Medvec A, et al. Clinical grade manufacturing of human alloantigen-  
628 reactive regulatory T cells for use in transplantation. *Am J Transplant* 2013; **13**(11): 3010-20.
- 629 11. Marin E, Bouchet-Delbos L, Renoult O, et al. Human tolerogenic dendritic  
630 cells regulate immune responses through lactate synthesis. *Cell Metab* 2019; **30**(6): 1075-90.
- 631 12. Hutchinson JA, Ahrens N, Geissler EK. MITAP-compliant characterization of human regulatory  
632 macrophages. *Transpl Int* 2017; **30**(8): 765-75.
- 633 13. Ekberg H, Tedesco-Silva H, Demirbas A, et al. Reduced exposure to calcineurin inhibitors in  
634 renal transplantation. *N Engl J Med* 2007; **357**(25): 2562-75.
- 635 14. Roufousse C, Simmonds N, Clahsen-van Groningen M, et al. A 2018 Reference Guide to the  
636 Banff Classification of Renal Allograft Pathology. *Transplantation* 2018; **102**(11): 1795-814.
- 637 15. Streitz M, Miloud T, Kapinsky M, et al. Standardization of whole blood immune phenotype  
638 monitoring for clinical trials: panels and methods from the ONE study. *Transplant Res* 2013; **2**(1): 17.
- 639 16. Kverneland AH, Streitz M, Geissler E, et al. Age and gender leucocytes variances and  
640 references values generated using the standardized ONE-Study protocol. *Cytometry A* 2016; **89**(6):  
641 543-64.
- 642 17. Fishman JA. Infection in Organ Transplantation. *Am J Transplant* 2017; **17**(4): 856-79.
- 643 18. Meijers RW, Litjens NH, de Wit EA, et al. Cytomegalovirus contributes partly to uraemia-  
644 associated premature immunological ageing of the T cell compartment. *Clin Exp Immunol* 2013;  
645 **174**(3): 424-32.
- 646 19. Meijers RW, Litjens NH, Hesselink DA, Langerak AW, Baan CC, Betjes MG. Primary  
647 Cytomegalovirus Infection Significantly Impacts Circulating T Cells in Kidney Transplant Recipients.  
648 *Am J Transplant* 2015; **15**(12): 3143-56.
- 649 20. Makwana N, Foley B, Fernandez S, et al. CMV drives the expansion of highly functional  
650 memory T cells expressing NK-cell receptors in renal transplant recipients. *Eur J Immunol* 2017; **47**(8):  
651 1324-34.
- 652 21. Ulrich C, Heine GH, Gerhart MK, Kohler H, Girndt M. Proinflammatory CD14+CD16+  
653 monocytes are associated with subclinical atherosclerosis in renal transplant patients. *Am J*  
654 *Transplant* 2008; **8**(1): 103-10.



- 655 22. Vereyken EJ, Kraaij MD, Baan CC, et al. A shift towards pro-inflammatory CD16+ monocyte  
656 subsets with preserved cytokine production potential after kidney transplantation. *PLoS One* 2013;  
657 **8**(7): e70152.
- 658 23. van den Bosch TPP, Hilbrands LB, Kraaijeveld R, et al. Pretransplant Numbers of CD16(+)  
659 Monocytes as a Novel Biomarker to Predict Acute Rejection After Kidney Transplantation: A Pilot  
660 Study. *Am J Transplant* 2017; **17**(10): 2659-67.
- 661 24. Braza F, Dugast E, Panov I, et al. Central Role of CD45RA- Foxp3hi Memory Regulatory T Cells  
662 in Clinical Kidney Transplantation Tolerance. *J Am Soc Nephrol* 2015; **26**(8): 1795-805.
- 663 25. Gray M, Gray D. Regulatory B cells mediate tolerance to apoptotic self in health: implications  
664 for disease. *Int Immunol* 2015; **27**(10): 505-11.
- 665 26. Appelgren D, Eriksson P, Ernerudh J, Segelmark M. Marginal-Zone B-Cells Are Main Producers  
666 of IgM in Humans, and Are Reduced in Patients With Autoimmune Vasculitis. *Front Immunol* 2018; **9**:  
667 2242.
- 668 27. Hutchinson JA, Brem-Exner BG, Riquelme P, et al. A cell-based approach to the minimization  
669 of immunosuppression in renal transplantation. *Transpl Int* 2008; **21**(8): 742-54.
- 670 28. Hutchinson JA, Riquelme P, Brem-Exner BG, et al. Transplant acceptance-inducing cells as an  
671 immune-conditioning therapy in renal transplantation. *Transpl Int* 2008; **21**(8): 728-41.
- 672 29. Riquelme P, Haarer J, Kammler A, et al. TIGIT+ iTregs elicited by human regulatory  
673 macrophages control T cell immunity. *Nat Comms* 2018; 9:2858 DOI:101038/s41467-018-05167-8.
- 674 30. Chandran S, Tang Q, Sarwal M, et al. Polyclonal Regulatory T Cell Therapy for Control of  
675 Inflammation in Kidney Transplants. *Am J Transplant* 2017; **17**(11): 2945-54.
- 676 31. Mathew JM, J HV, LeFever A, et al. A Phase I Clinical Trial with Ex Vivo Expanded Recipient  
677 Regulatory T cells in Living Donor Kidney Transplants. *Scientific Rep* 2018; **8**(1): 7428.
- 678 32. Todo S, Yamashita K, Goto R, et al. A pilot study of operational tolerance with a regulatory T-  
679 cell-based cell therapy in living donor liver transplantation. *Hepatology* 2016; **64**(2): 632-43.

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**Fig. 1:** ONE Study design and patient disposition for the multicenter RGT and six monocenter CTG trials. RGT = Reference Group Trial; CTG = Cell Therapy Group trials; Mreg: regulatory macrophages; ATDC: autologous tolerogenic dendritic cells; pTreg-1 / pTreg-2: polyclonal regulatory T cells; darTreg-sBC: donor-alloantigen reactive Treg; darTreg-CSB: costimulatory blockade generated Treg.

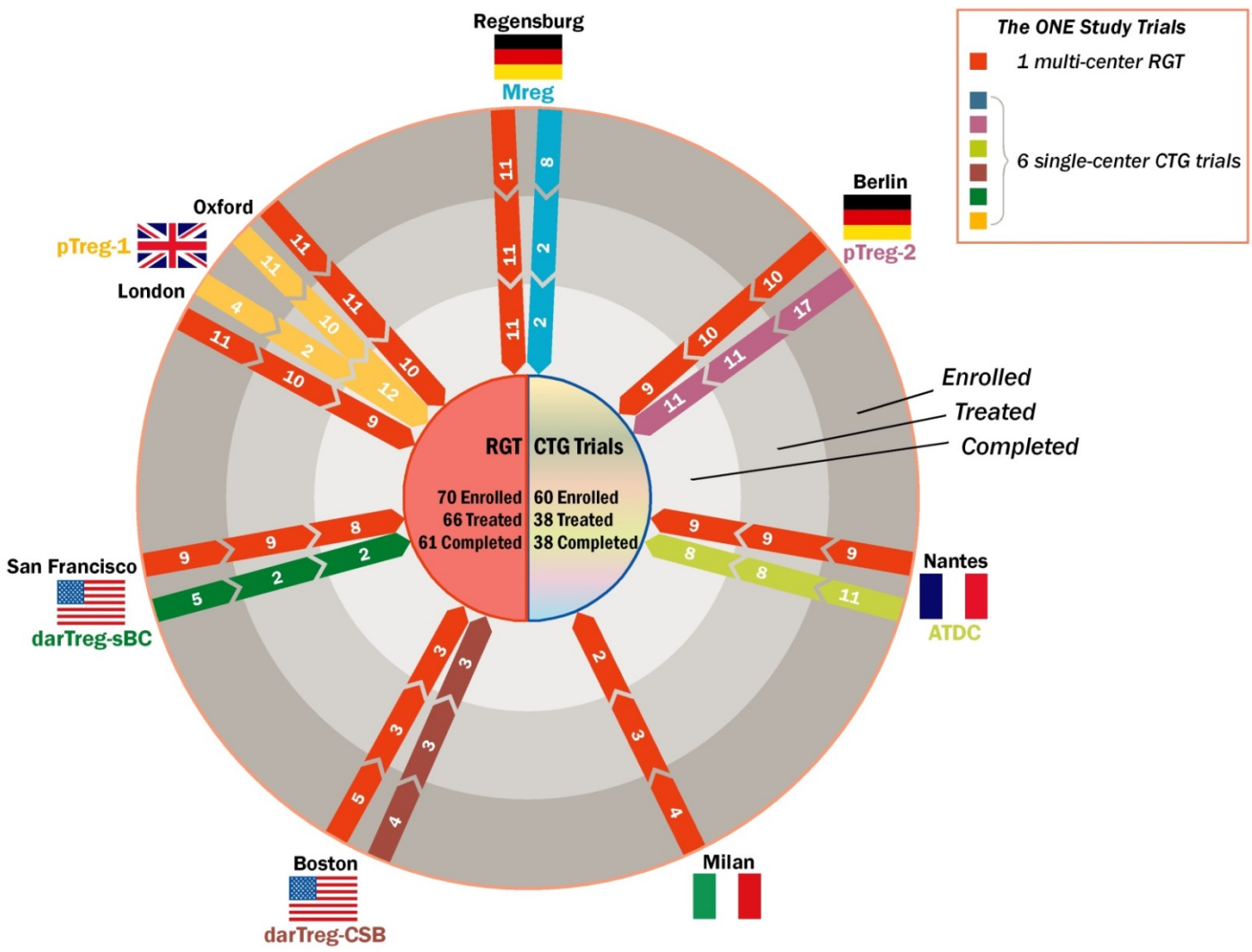
**Fig. 2:** Primary endpoint (BCAR) data. 2A). Kaplan-Meier estimates of the cumulative BCAR-free survival probability in the RGT (N=66) and CTG (N=38) intention-to-treat analysis sets (87.7 % vs. 84.2 % at 60 weeks). Censored patients marked with ticks. 2B). Severity of first BCAR episode by central pathological diagnosis and response to treatment. \* One patient treated with low-dose oral steroids and by not tapering immunosuppression. 2C). Kaplan-Meier estimates of the cumulative BCAR-free survival probability in the RGT (N=47) and CTG (N=32) per-protocol analysis sets (82.8 % vs. 81.3 % at 60 weeks). Censored patients marked with ticks. RGT = Reference Group Trial; CTG = Cell Therapy Group trials; BCAR = biopsy-confirmed acute rejection; TCMR = T cell-mediated rejection; ABMR = antibody-mediated rejection.

**Fig. 3:** ONE Study safety data (normalized). 3A) Incidence rate of treatment-emergent SAEs by MedDRA primary SOC. 3B) Incidence rate of treatment-emergent infections (all AEs) by study site. 3C) Incidence proportion of treatment-emergent infections (all AEs) over time. 3D) Incidence rate of treatment-emergent infections (all AEs) by MedDRA HLGT. 3E) Incidence rate of treatment-emergent viral infections (all AEs) by MedDRA HLT. All adverse events coded using MedDRA version 20.1. Treatment-emergent (S)AEs are events with onset date equal to or after first dose of any study drug. All events coded to the MedDRA PT: "Transplant rejection" are excluded, since rejection was measured as the primary efficacy endpoint. RGT = Reference Group Trial; CTG = Cell Therapy Group trials; SOC = System Organ Class; HLGT = High Level Group Term; HLT = High Level Term; PSY = Patient study years; NEC = Not elsewhere classified.

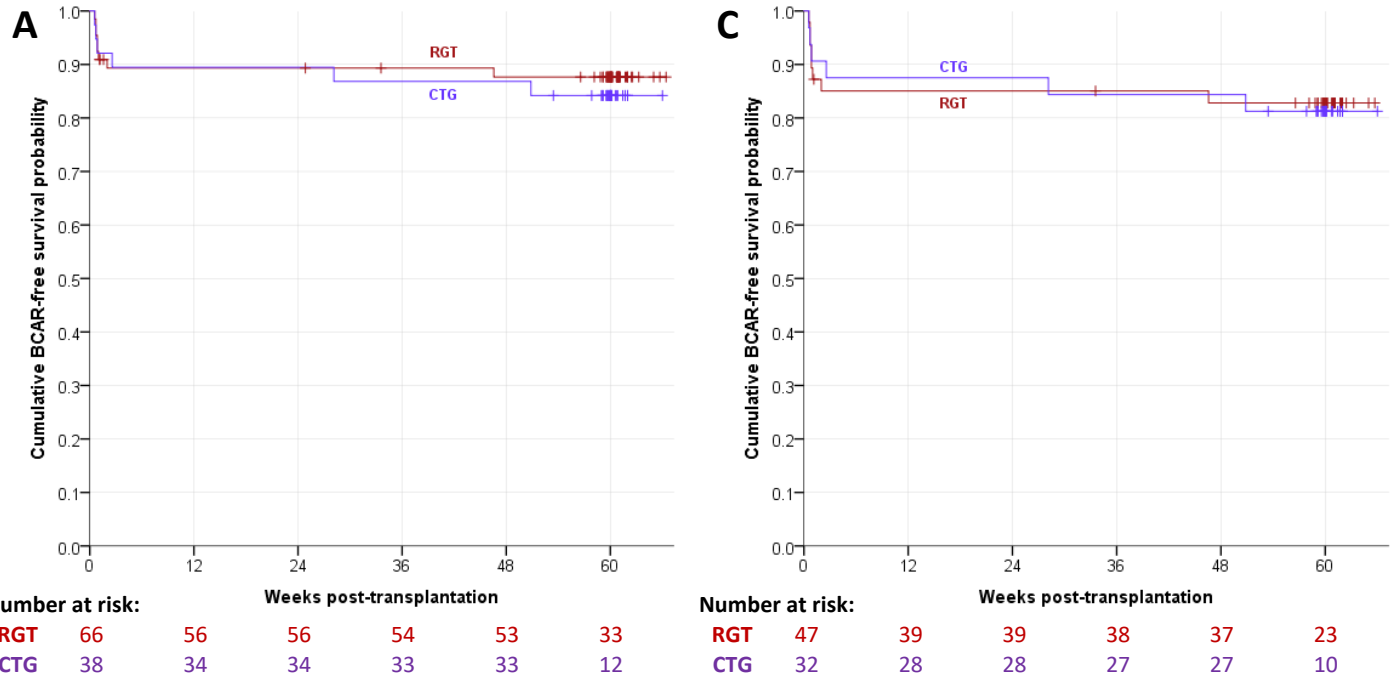
**Fig. 4:** Leukocyte subset alterations in ESRD patients and time-dependent changes after kidney transplantation. A) Principal component analysis revealing the differences in leukocyte subset between whole blood samples from end stage renal renal disease (ESRD, n= 70) and healthy controls (HC, n= 98). B) Box-and-whiskers plots of absolute numbers from leukocyte subpopulations with highest influence at the PCA shown in A. C) Time-dependent changes from visit 1 prior to transplantation (V01) to visit 10 at 60 weeks post-transplant (V10) of monocyte, B cell, CD4<sup>+</sup> and CD8<sup>+</sup> T cell subset composition (stacked bars of mean proportions) in whole blood samples of RGT patients (n=59). Statistical analysis by Kruskal-Wallis-Test. \* p<0.05, \*\* p<0.01

**Fig. 5:** Differences in post-transplant changes between RGT and CTG patients. A) Differences in post-transplant changes in regulatory T cells. Box and whisker plots of absolute numbers and proportions of CD4<sup>+</sup>CD25<sup>high</sup>CD127<sup>low</sup> Tregs as well as % CD4<sup>+</sup> T cells with demethylated TSDR in whole blood samples collected pre-transplant (V01) and at the end of the observation period (15 months post-transplant, V10) from RGT (n=59) and CTG patients (n=38) measured as described in material and methods. B) Differences in post-transplant changes in CD8<sup>+</sup> T cell subpopulations. Box and whisker plots of absolute numbers of CD8<sup>+</sup>CD28<sup>+</sup>, CD8<sup>+</sup>CD45RA<sup>+</sup>CCR7<sup>-</sup> T<sub>EMRA</sub> and CD8<sup>+</sup>CD57<sup>+</sup> chronically activated cells in whole blood samples collected pre-transplant (V01) and at the end of the observation period (15 months post-transplant, V10) from RGT (n=59) and CTG patients (n=38). C) Differences in post-transplant changes in marginal zone-like B cells and dendritic cell subpopulation. Box and whisker plots of absolute numbers and proportions of marginal zone-like B cells, CD16<sup>+</sup> mDCs and pDCs in whole blood samples collected pre-transplant (V01) and at the end of the observation period (15 months post-transplant, V10) from RGT (n=59) and CTG patients (n=38). Statistical analysis by Wilcoxon matched-pairs signed rank and Dunn's multiple comparison test. \* p<0.05, \*\* p<0.01, \*\*\* p<0.001, \*\*\*\* p<0.0001

**Figure 1**



# Figure 2



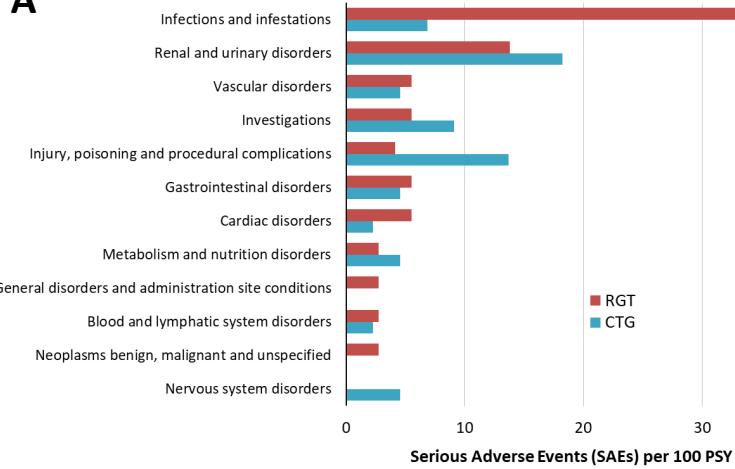
**B**

Severity of 1st BCAR episode

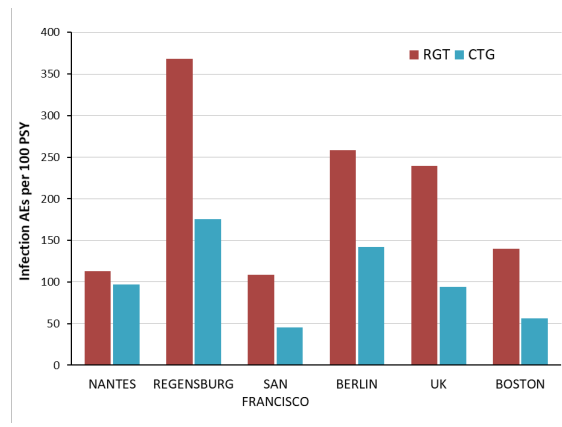
	RGT (N=8)	CTG (N=6)
<b>Central pathological diagnosis</b>		
Acute TCMR IA	1 (12.5 %)	1 (16.7 %)
Acute TCMR IIA	3 (37.5 %)	2 (33.3 %)
Acute TCMR IB	1 (12.5 %)	1 (16.7 %)
Acute TCMR IIB	0 (0.0 %)	2 (33.3 %)
Borderline changes	3 (37.5 %)	0 (0.0 %)
<b>ABMR diagnosed locally?</b>		
Yes	1 (12.5 %)	2 (33.3 %)
No	7 (87.5 %)	4 (66.7 %)
<b>Response to treatment</b>		
Glucocorticoid-responsive	4 (50.0 %)	3 (50.0 %)
Responsive to depleting antibody treatment	3 (37.5 %)	3 (50.0 %)
Not applicable*	1 (12.5 %)	0 (0.0 %)

**Figure 3**

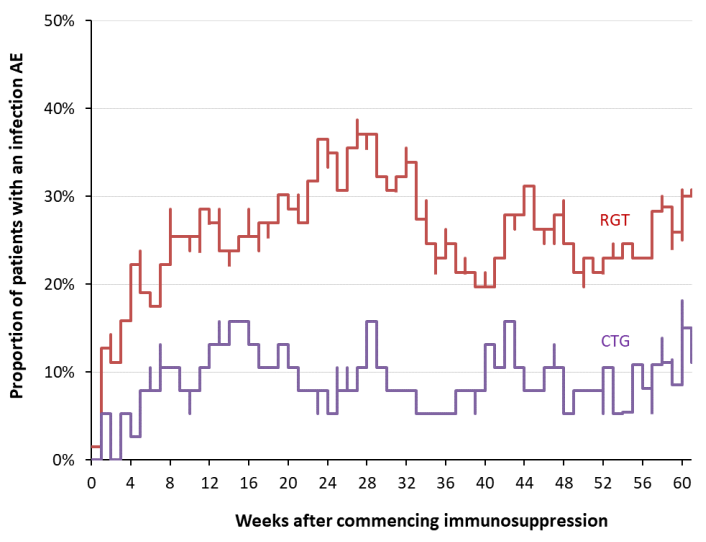
**A**



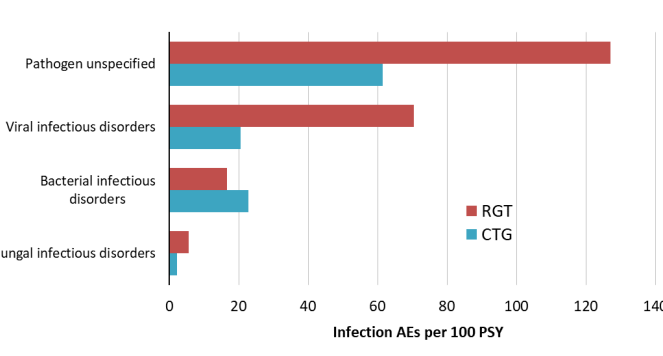
**B**



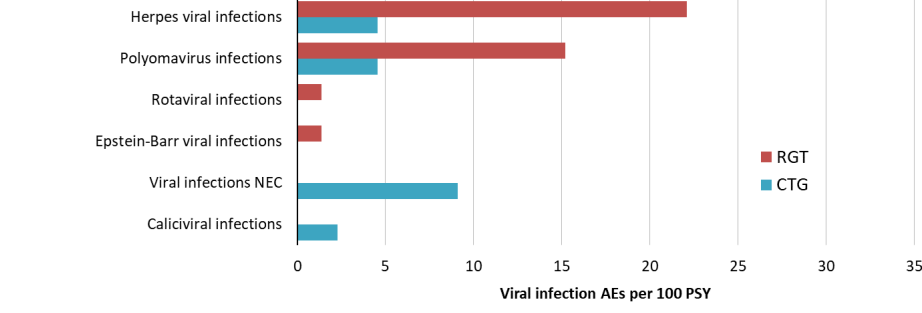
**C**



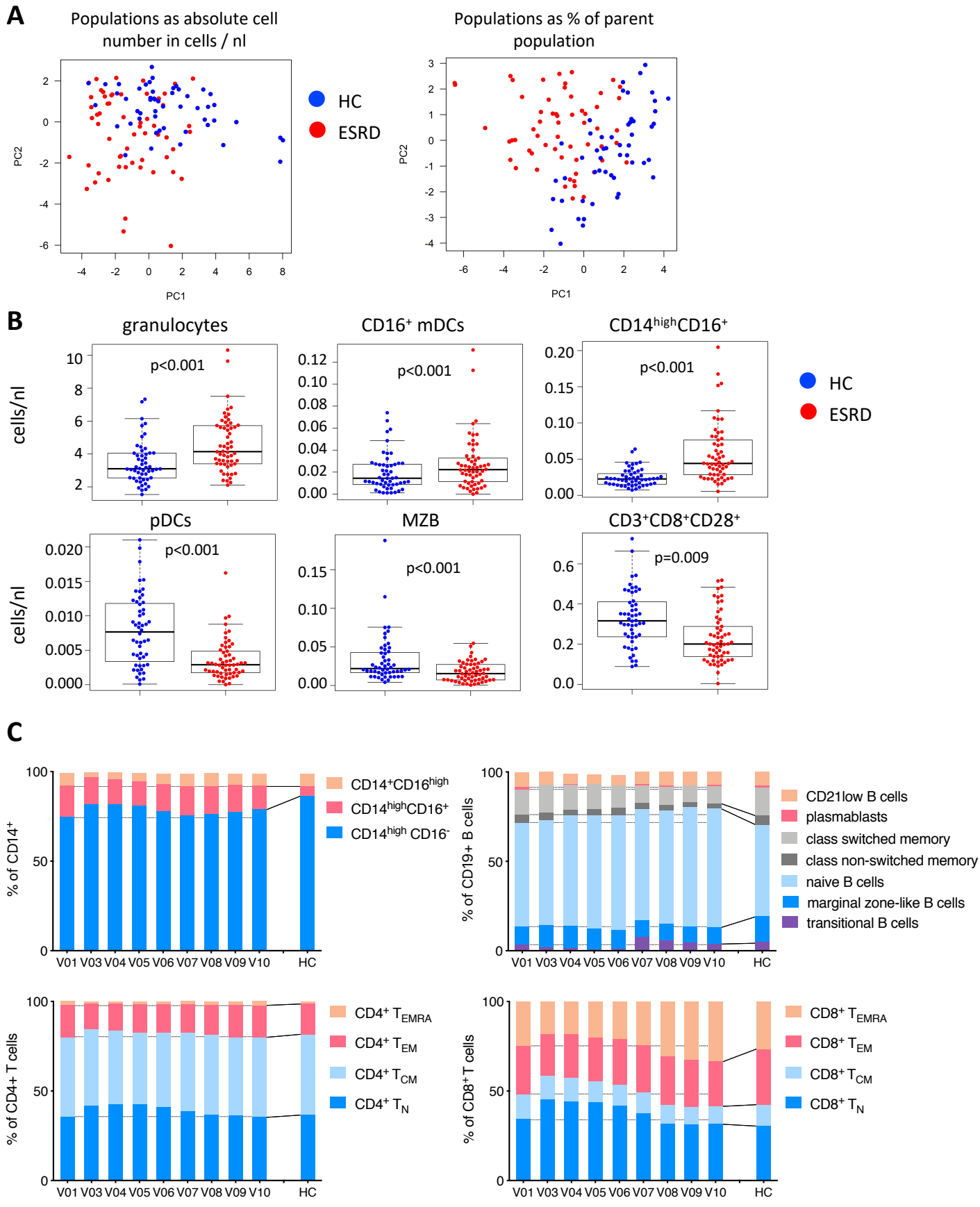
**D**



**E**

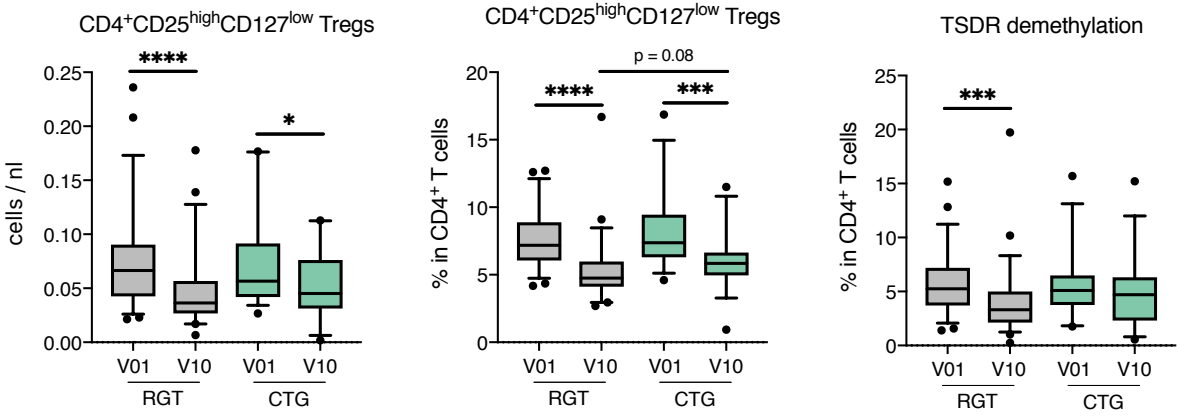


**Figure 4**

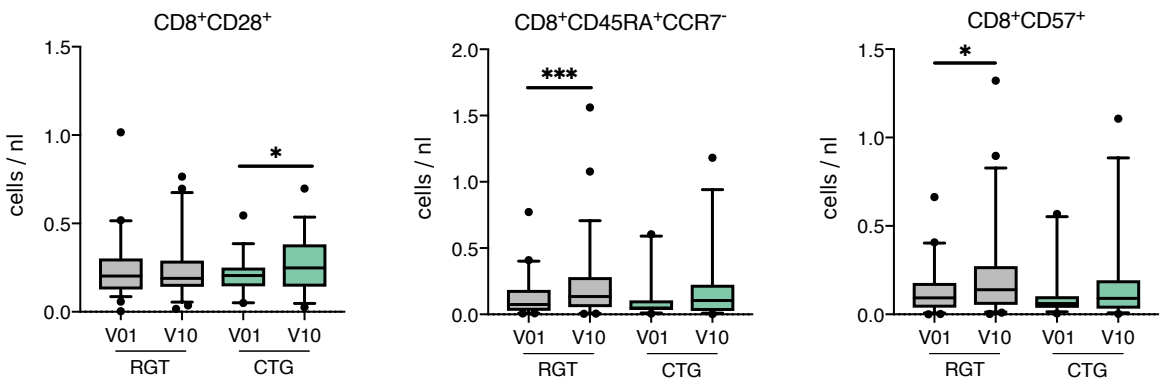


**Figure 5**

**A Regulatory T cells**



**B CD8<sup>+</sup> T cell subpopulations**



**C Marginal zone-like B cells and dendritic cell subpopulations**

