

1 **Additional file 1**

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3 **Table S1: Linear regression analysis for percentages of CD4⁺T_{EM} cells, T_{EM}1,**

4 **and T_{EM}17 cells between r-GPA patients and HCs**

	Predicting variable	B	(95% CI)	p-value
CD4⁺T_{EM} cells	Group (r-GPA and HCs)	17.3	(12,104 – 22,516)	<0.0001
	CMV and age	17.1	(12,161 – 21,969)	<0.0001
Log(T_{EM}1) cells	Group (r-GPA and HCs)	-.121	(-.219 – -.023)	0.016
	CMV and age	-.107	(-.198 – -.017)	0.021
T_{EM}17 cells	Group (r-GPA and HCs)	5,73	(2,321 – 9,132)	0.001
	CMV and age	5,29	(2,235 – 8,339)	0.001
Log(T_{EM}1) cells	Group (relapse and non-relapse r-GPA patients)	-.145	(-.288 – -.002)	0.047
	CMV and age	-.135	(-.263 – -.007)	0.040
T_{EM}17 cells	Group (relapse and non-relapse r-GPA patients)	4.80	(-.227 – 9.823)	0.061
	CMV and age	4.55	(.145 – 8.96)	0.043

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