Correlations: Benchmarks and Human-Model Alignment 0.67 0.43 0.03 0.50 0.68 0.64 0.67 0.44 ifeval 1.00

1.0

-0.8

- 0.6 **└**

-0.2

bbh 0.44 1.00 0.72 0.88 0.50 0.94 0.74 0.70 0.64 0.72 1.00 0.89 0.19 0.84 0.61 0.57 0.51 0.67 math 0.37 0.43 0.88 0.89 1.00 0.88 0.56 0.51 0.46 gpga 0.03 0.50 0.19 0.37 1.00 0.45 0.36 0.34 0.33 musr 0.50 0.94 0.84 0.88 0.45 1.00 0.71 0.68 0.62 mmlu 0.74 0.61 0.56 0.36 0.71 1.00 0.99 0.97 Procrustes R2 0.68

0.64 0.70 0.57 0.51 0.34 0.68 0.99 1.00 0.98 cka 0.67 0.64 0.51 0.46 0.33 0.62 0.97 0.98 1.00 **RSM**