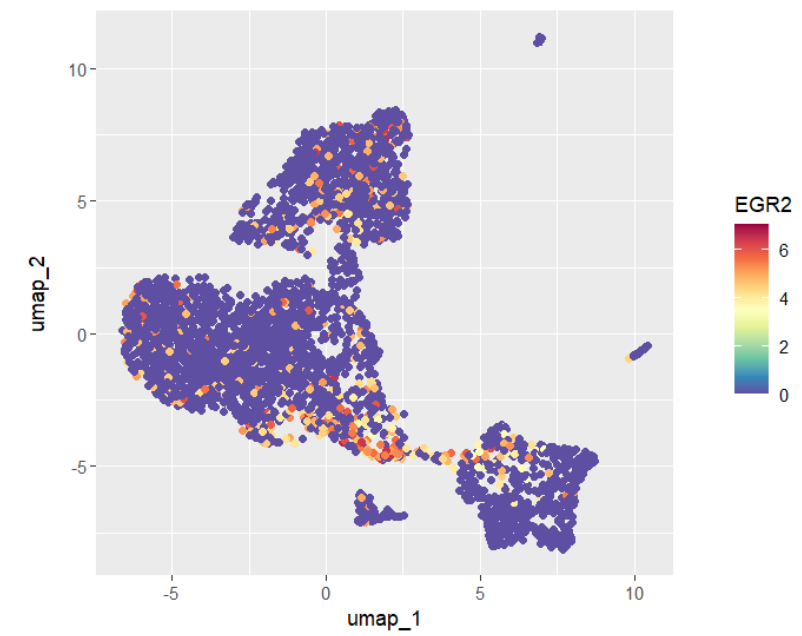
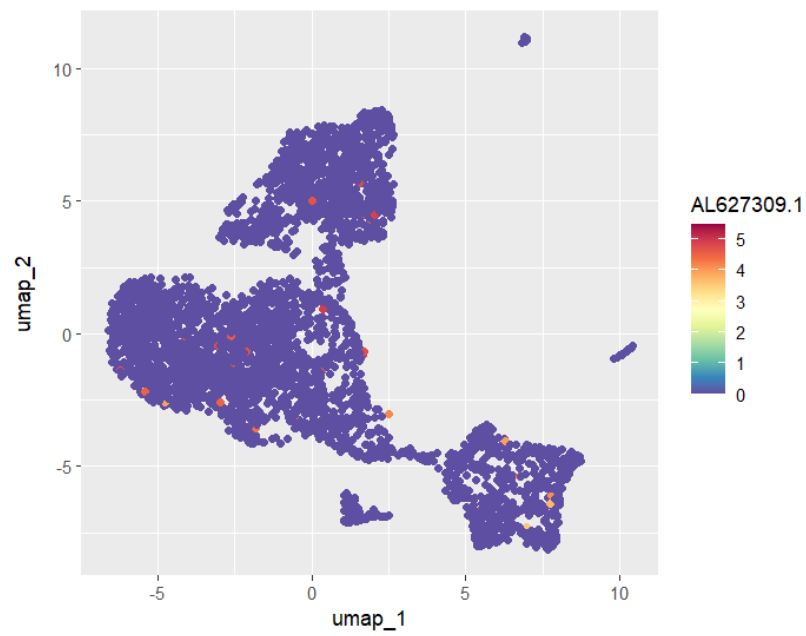
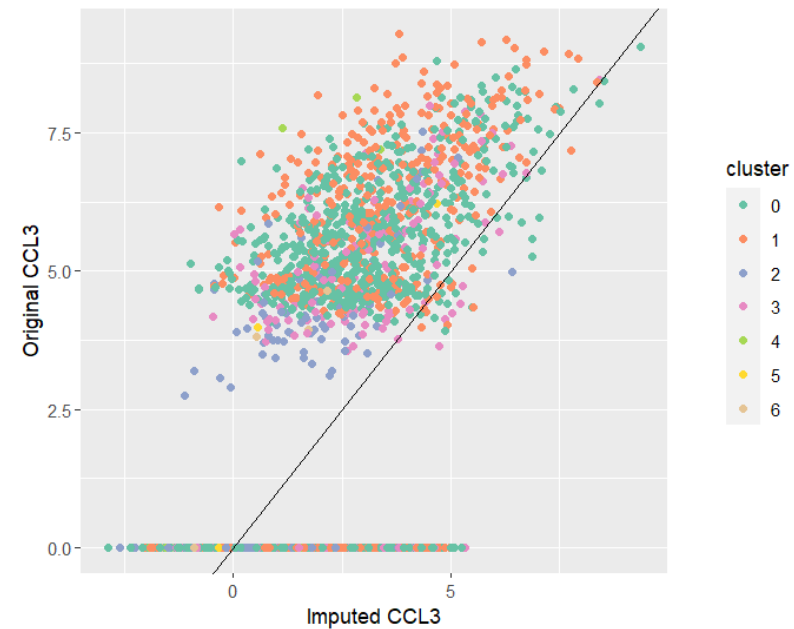
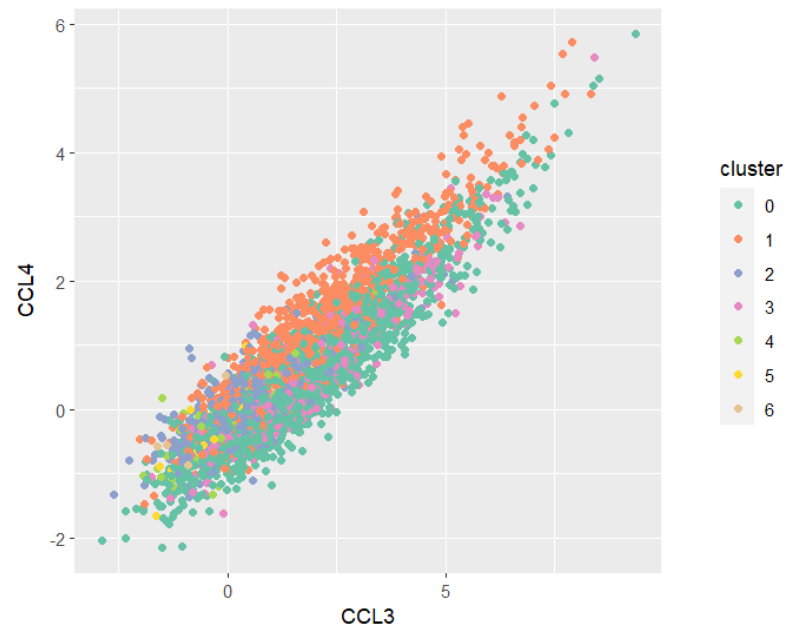
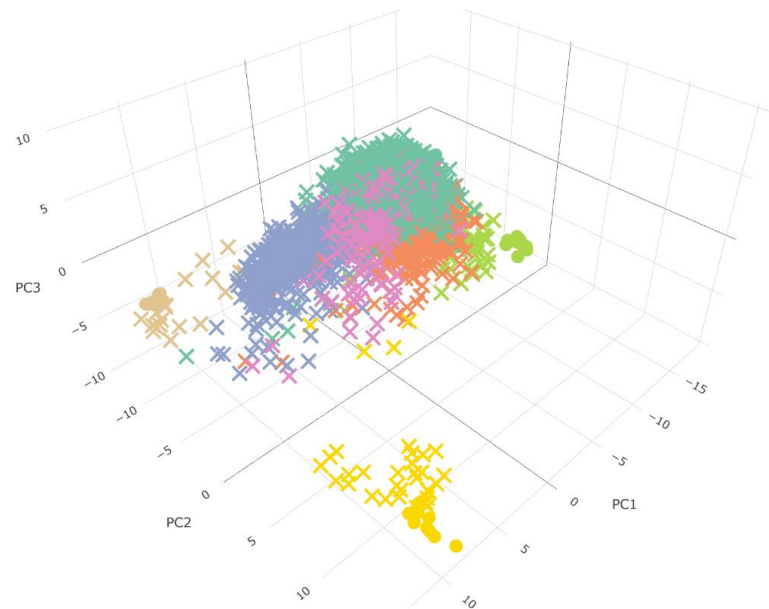
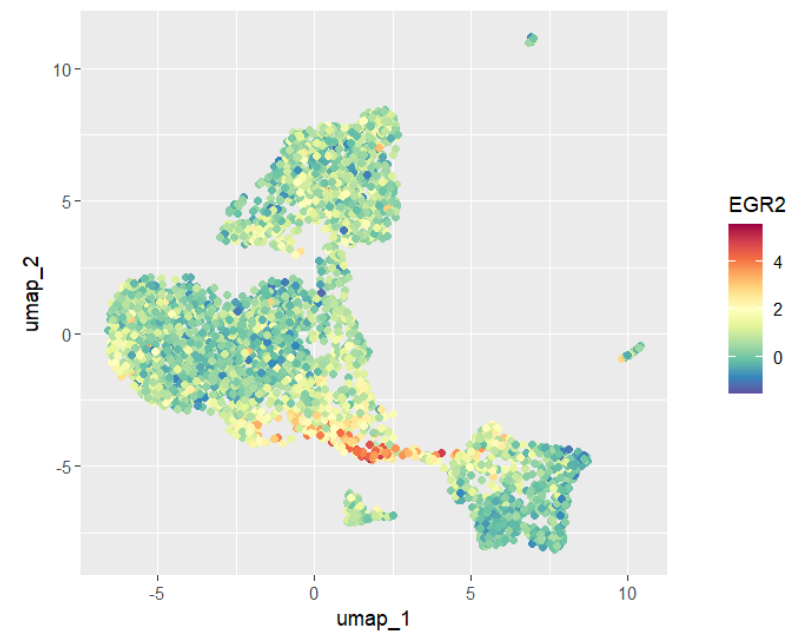
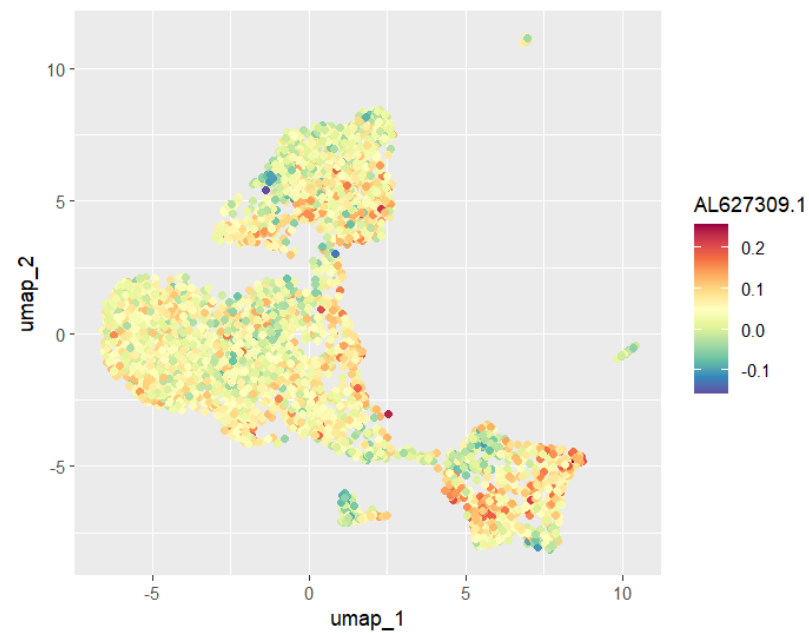


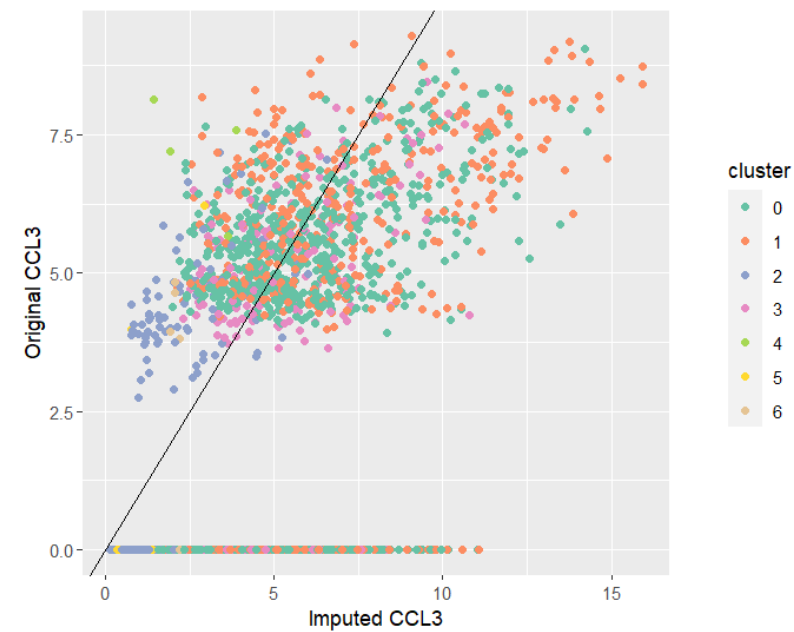
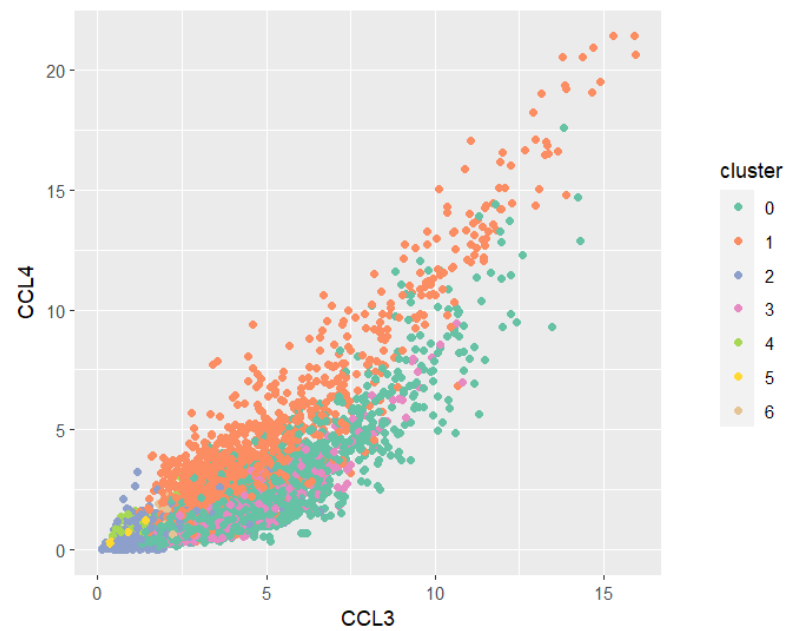
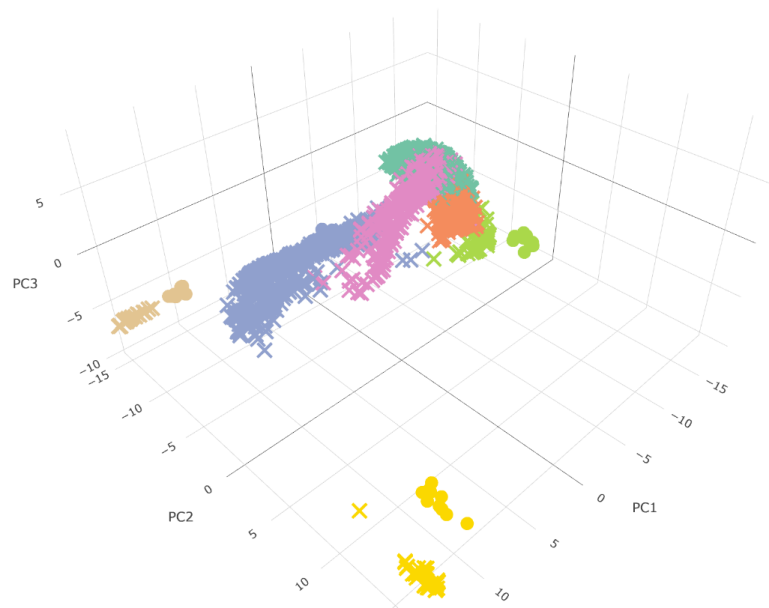
Original



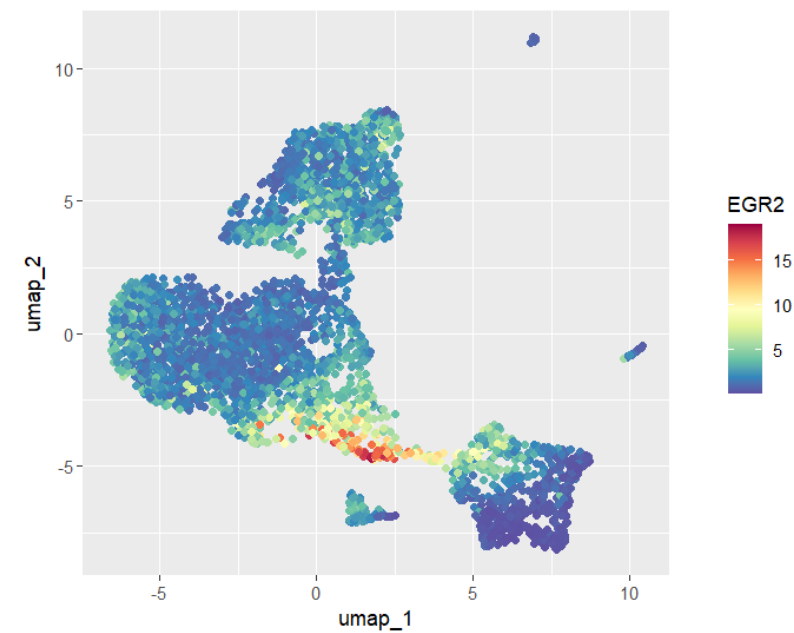
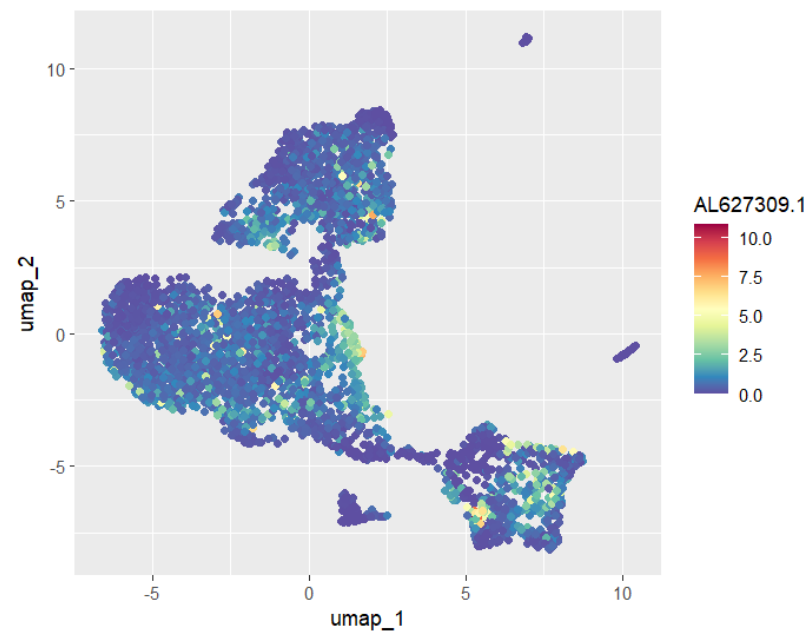


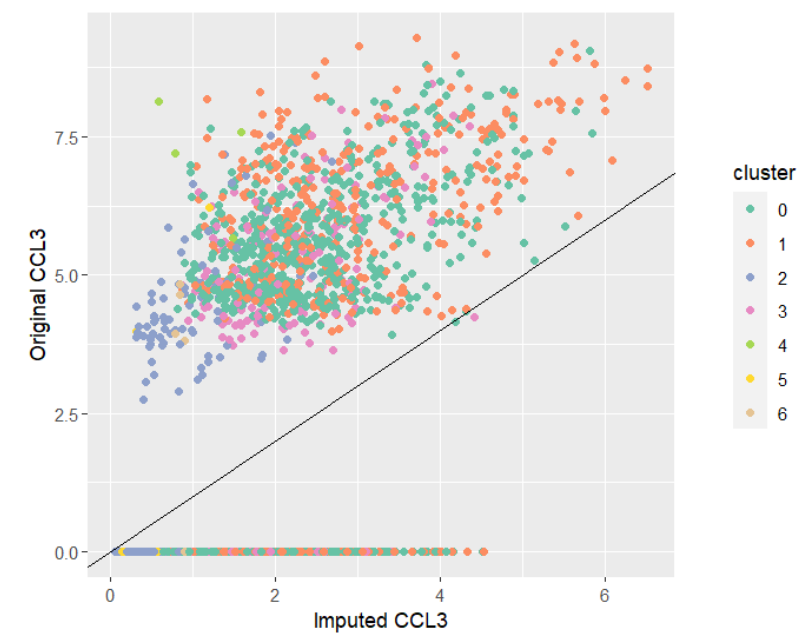
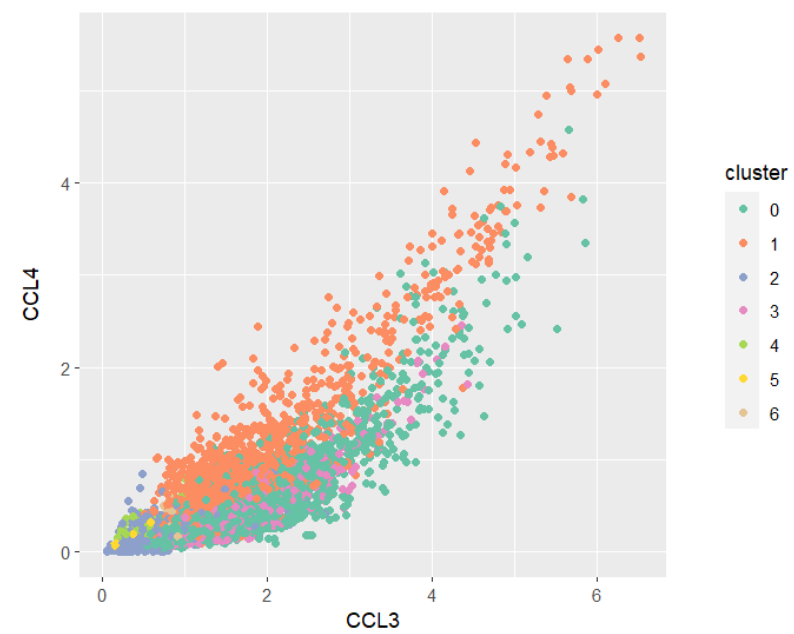
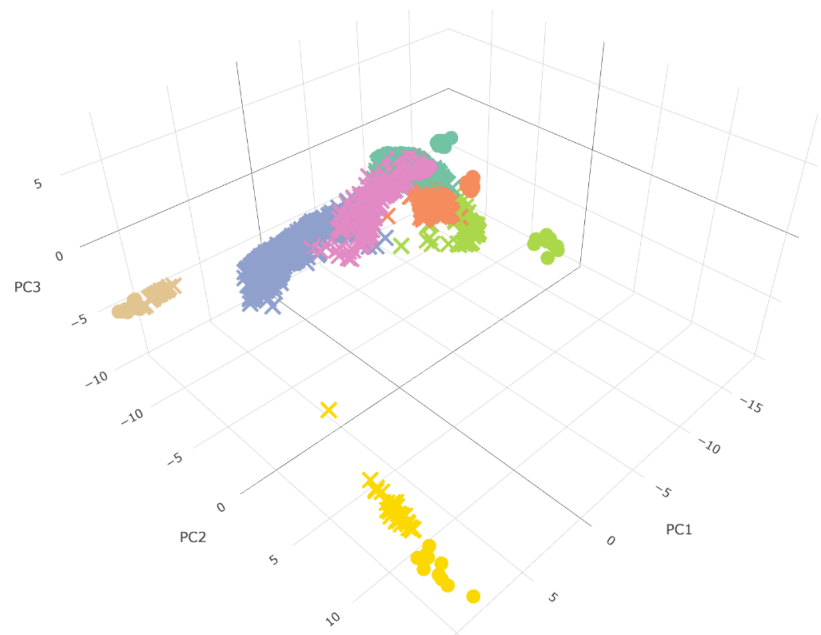
Low rank approximation



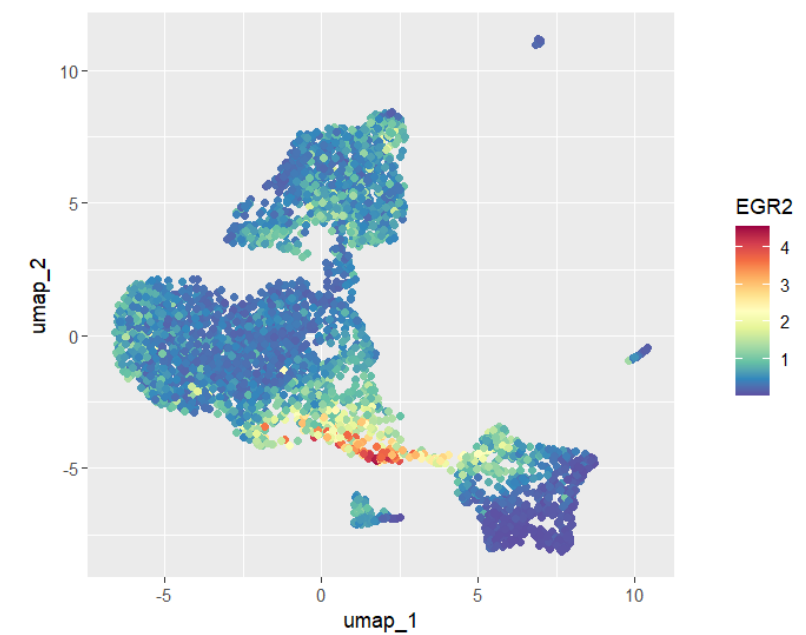
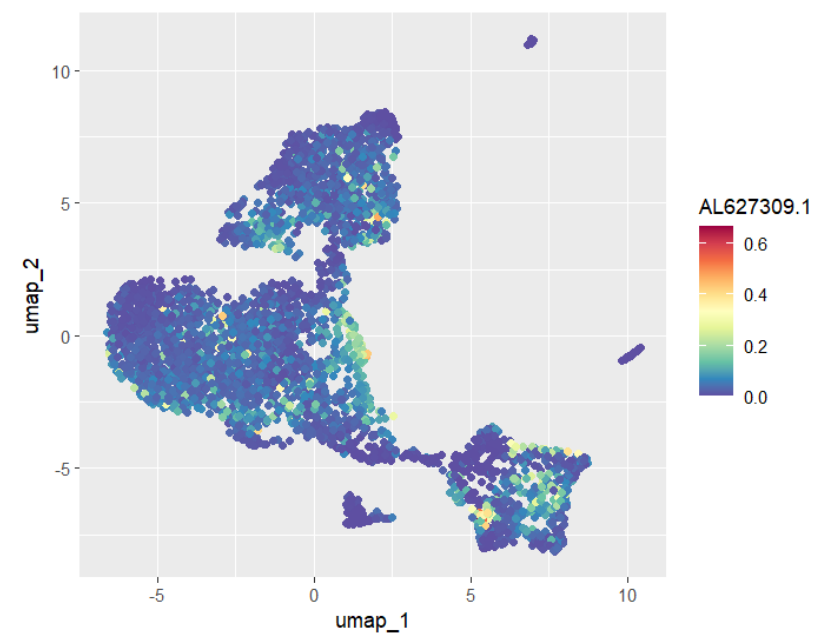


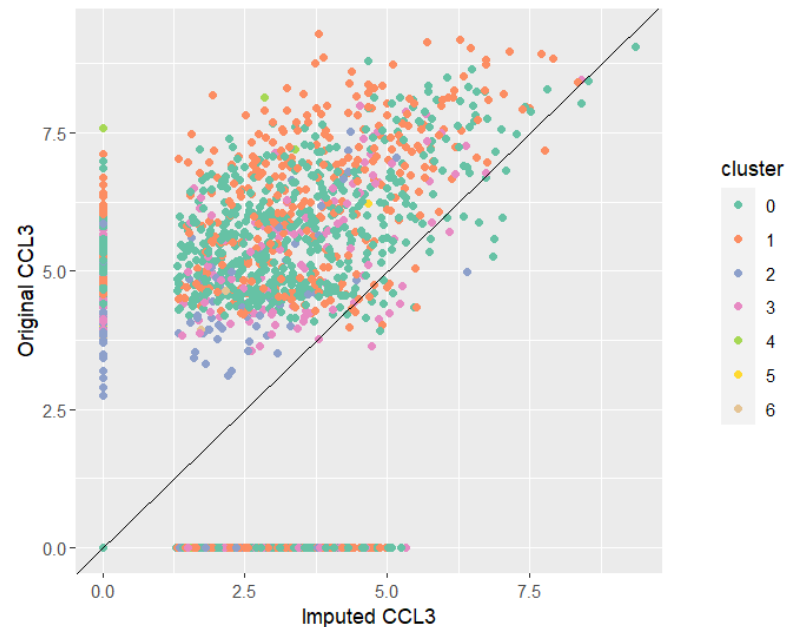
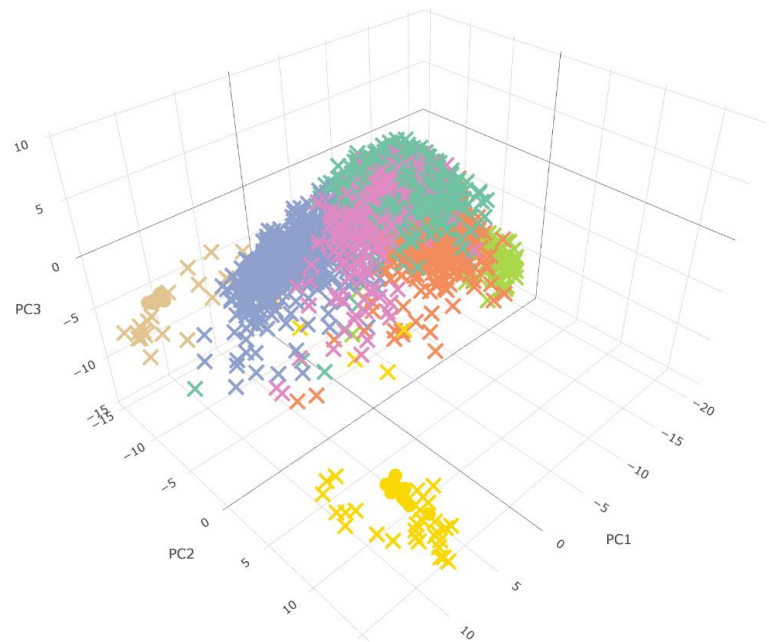
Sincast + Median matching





Sincast





Zero preserving Low rank approximation

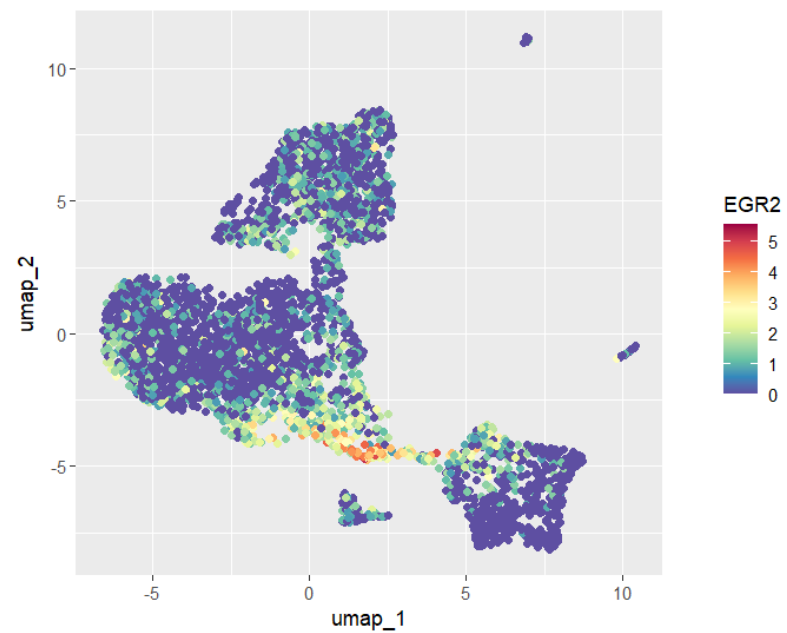
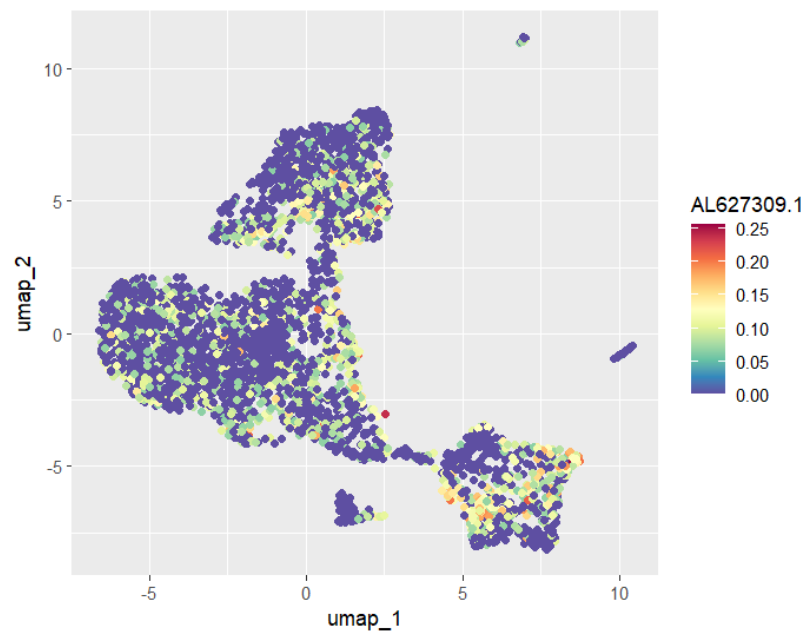
Article | [Open access](#) | Published: 11 January 2022

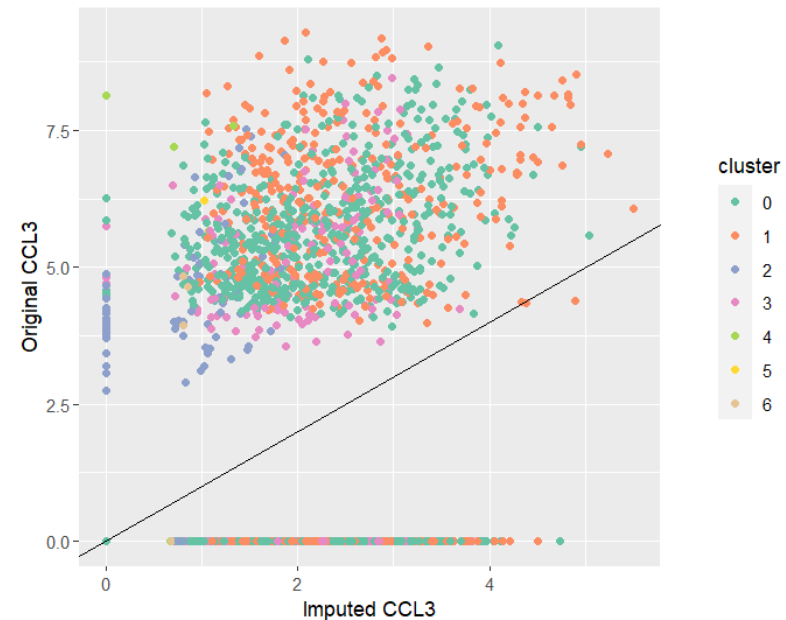
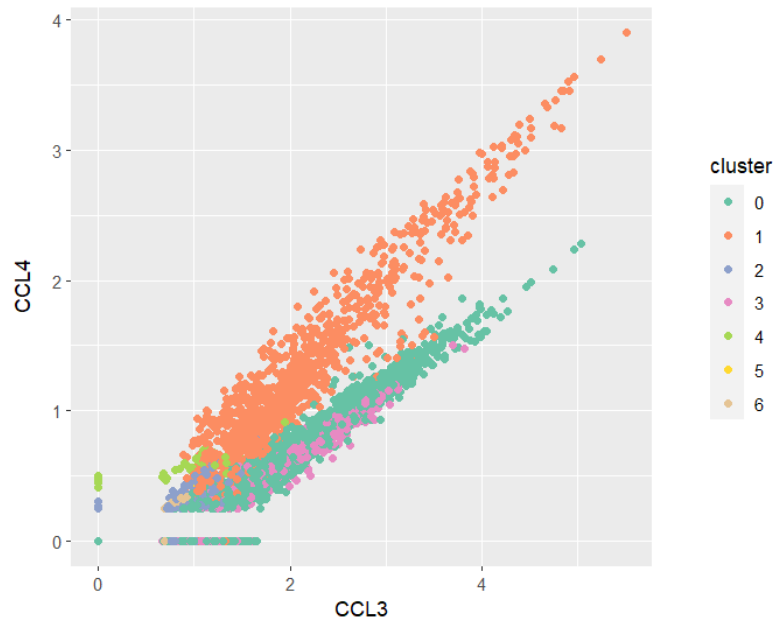
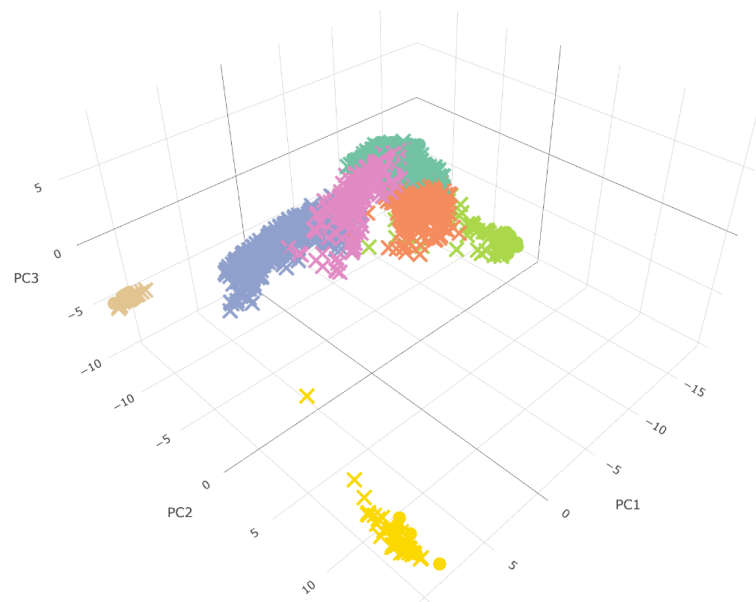
Zero-preserving imputation of single-cell RNA-seq data

[George C. Linderman](#), [Jun Zhao](#), [Manolis Roulis](#), [Piotr Bielecki](#), [Richard A. Flavell](#), [Boaz Nadler](#) & [Yuval Kluger](#) 

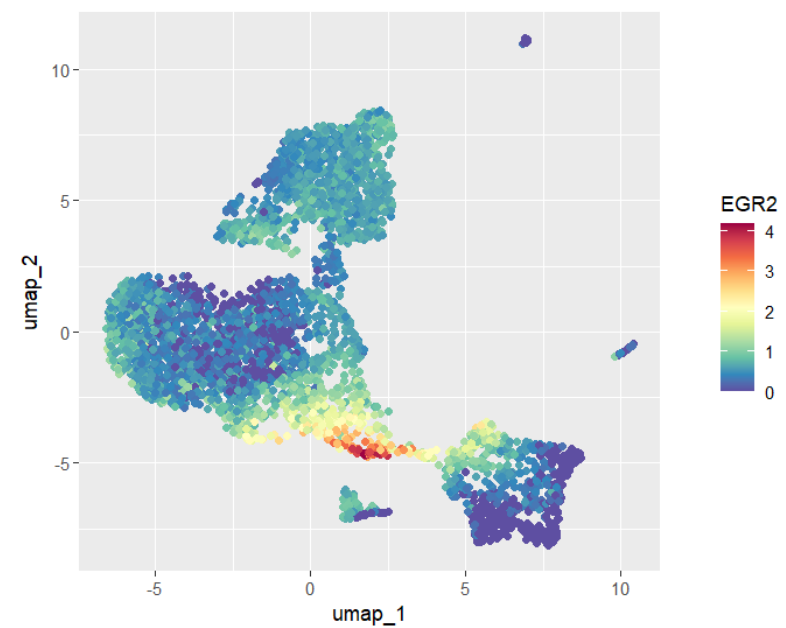
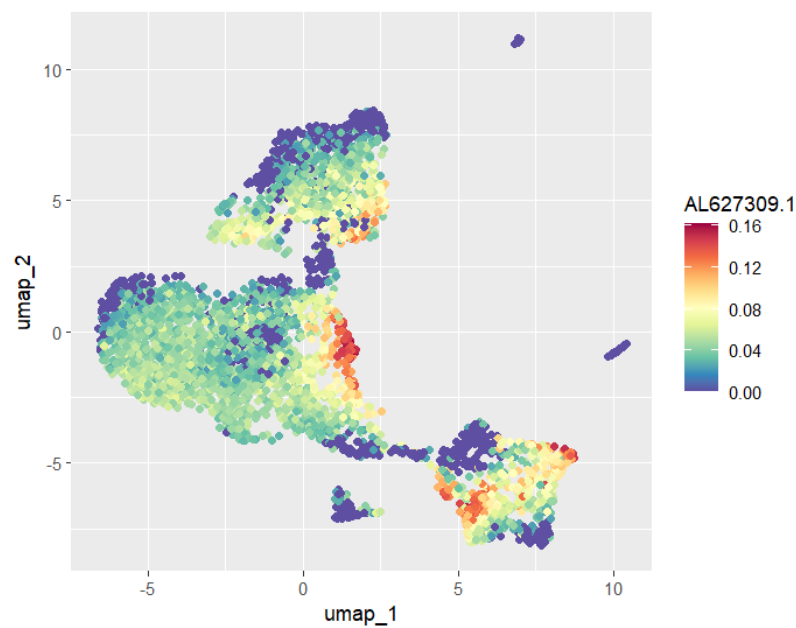
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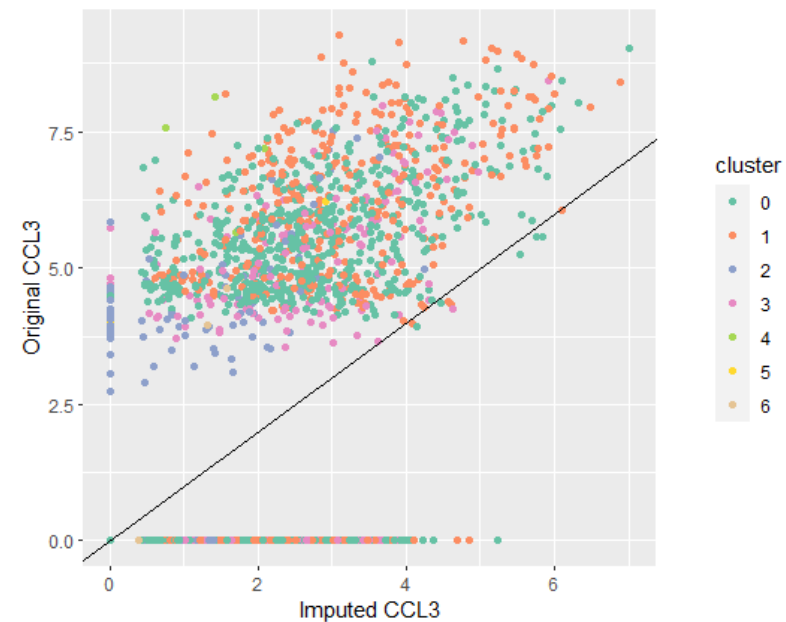
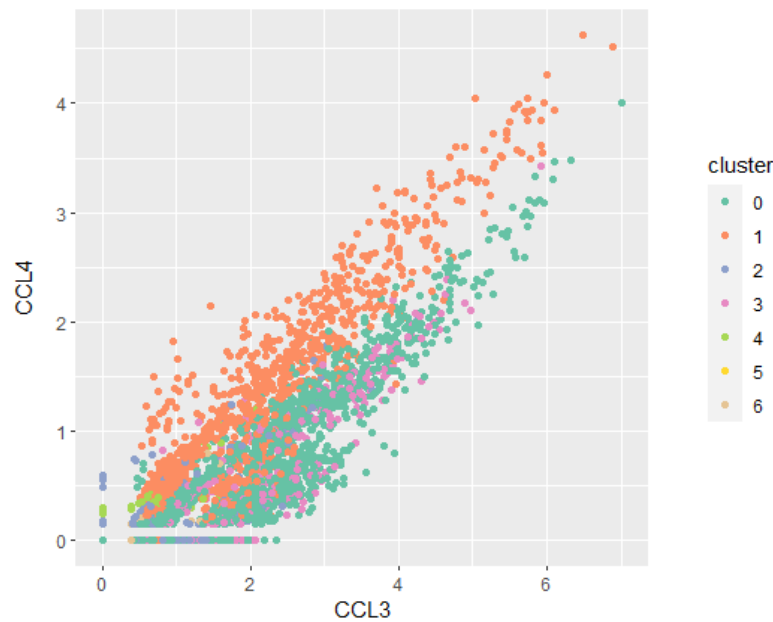
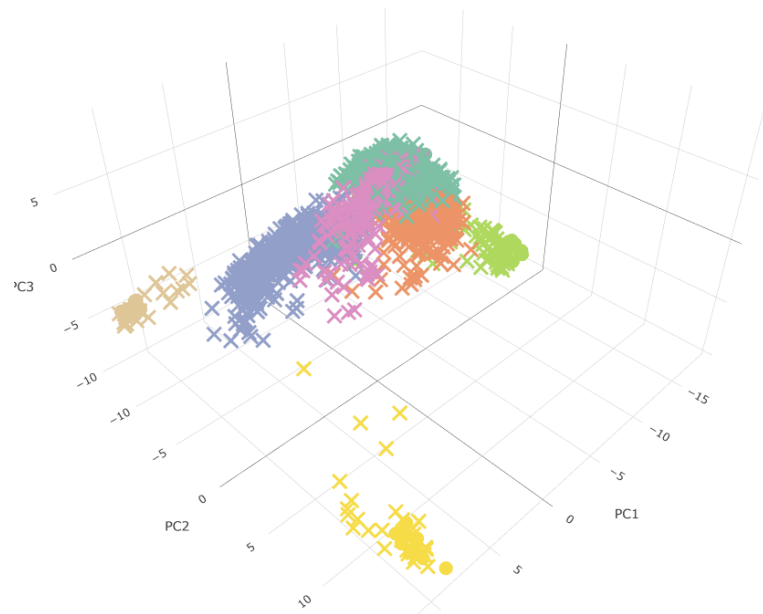
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Zero preserving
Low rank approximation on Sincast





Final Sincast

