

1.2 Release Notes

T	Key	Summary	Status
	BARNA-257	Error in read ID: could not parse read identifier in indexed bam generated by GEMtools	RESOLVED
	BARNA-256	When an error occurs solving the linear system, the matrix is written on stderr and an exception is thrown even if the deconvolution continues	RESOLVED
	BARNA-248	Capacitor complains about READ_DESCRIPTOR when given a unindexed BAM file	RESOLVED
	BARNA-247	Capacitor still wants the -p <parameterfile> option even if I specify all mandatory paramters on the command line	RESOLVED
	BARNA-246	UnixSorter line interceptor called for each merge step instead of just the final one	RESOLVED
	BARNA-242	When counting Splice Junctions and Introns, output the correct GeneID of a locus if multiple Genes are contained	RESOLVED
	BARNA-240	Capacitor runs out of Heap Space with some BAM files	RESOLVED
	BARNA-239	Parameter Schema relative path parser extends Windows paths	RESOLVED
	BARNA-238	Wrong number of reads detected in some BAM input files	RESOLVED

9 issues