Expression Noise (CV) by Bewick Group in Lineages (Mean > 0.5 Filtered)

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Statistical Comparisons for Expression Noise (CV) by Bewick Group in Lineages (Mean > 0.5 Filtered)

Context	Filter_Tag	Specific_Context	Group_Type	Comparison	P_value_Formatted
CV Expr by Lineage	Mean > 0.5 Filtered	rootcap	bewick_group	Overall (Kruskal–Wallis)	9.2644e–03
CV Expr by Lineage	Mean > 0.5 Filtered	rootcap	bewick_group	gbM vs Unmethylated	9.2668e–03
CV Expr by Lineage	Mean > 0.5 Filtered	epidermis	bewick_group	Overall (Kruskal–Wallis)	3.9916e–04
CV Expr by Lineage	Mean > 0.5 Filtered	epidermis	bewick_group	gbM vs Unmethylated	3.9930e–04
CV Expr by Lineage	Mean > 0.5 Filtered	cortex	bewick_group	Overall (Kruskal–Wallis)	7.8546e–03
CV Expr by Lineage	Mean > 0.5 Filtered	cortex	bewick_group	gbM vs Unmethylated	7.8567e–03
CV Expr by Lineage	Mean > 0.5 Filtered	xpp	bewick_group	Overall (Kruskal–Wallis)	9.2741e–01
CV Expr by Lineage	Mean > 0.5 Filtered	xpp	bewick_group	gbM vs Unmethylated	9.2749e–01
CV Expr by Lineage	Mean > 0.5 Filtered	endodermis	bewick_group	Overall (Kruskal–Wallis)	3.0397e–01
CV Expr by Lineage	Mean > 0.5 Filtered	endodermis	bewick_group	gbM vs Unmethylated	3.0401e–01
CV Expr by Lineage	Mean > 0.5 Filtered	ppp	bewick_group	Overall (Kruskal–Wallis)	6.0794e–02
CV Expr by Lineage	Mean > 0.5 Filtered	ppp	bewick_group	gbM vs Unmethylated	6.0806e–02
CV Expr by Lineage	Mean > 0.5 Filtered	procambium	bewick_group	Overall (Kruskal–Wallis)	7.0309e–01
CV Expr by Lineage	Mean > 0.5 Filtered	procambium	bewick_group	gbM vs Unmethylated	7.0315e–01
CV Expr by Lineage	Mean > 0.5 Filtered	phloem	bewick_group	Overall (Kruskal–Wallis)	3.4203e–01
CV Expr by Lineage	Mean > 0.5 Filtered	phloem	bewick_group	gbM vs Unmethylated	3.4208e–01
CV Expr by Lineage	Mean > 0.5 Filtered	xylem	bewick_group	Overall (Kruskal–Wallis)	3.7436e–05
CV Expr by Lineage	Mean > 0.5 Filtered	xylem	bewick_group	gbM vs Unmethylated	3.7451e–05