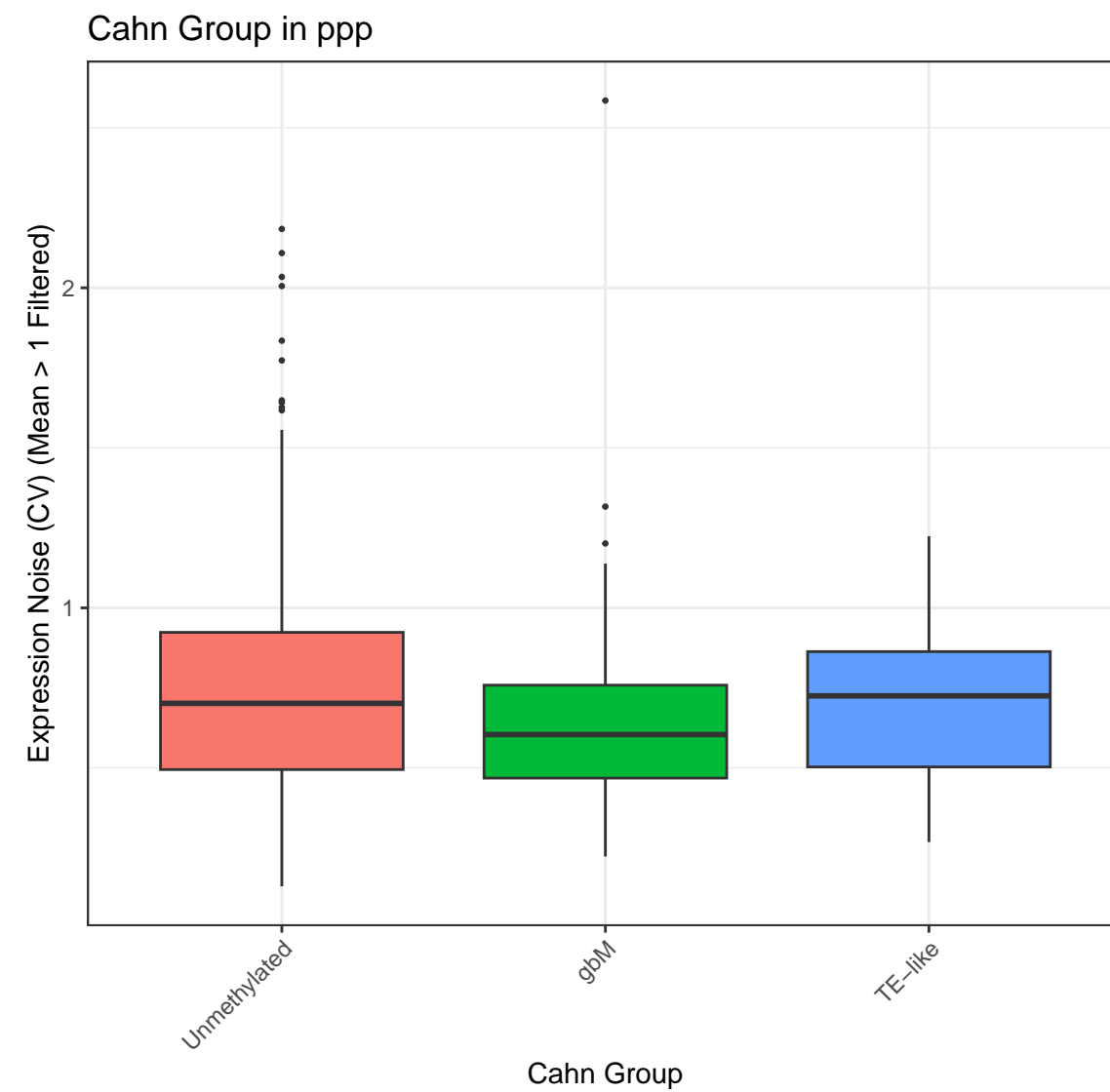
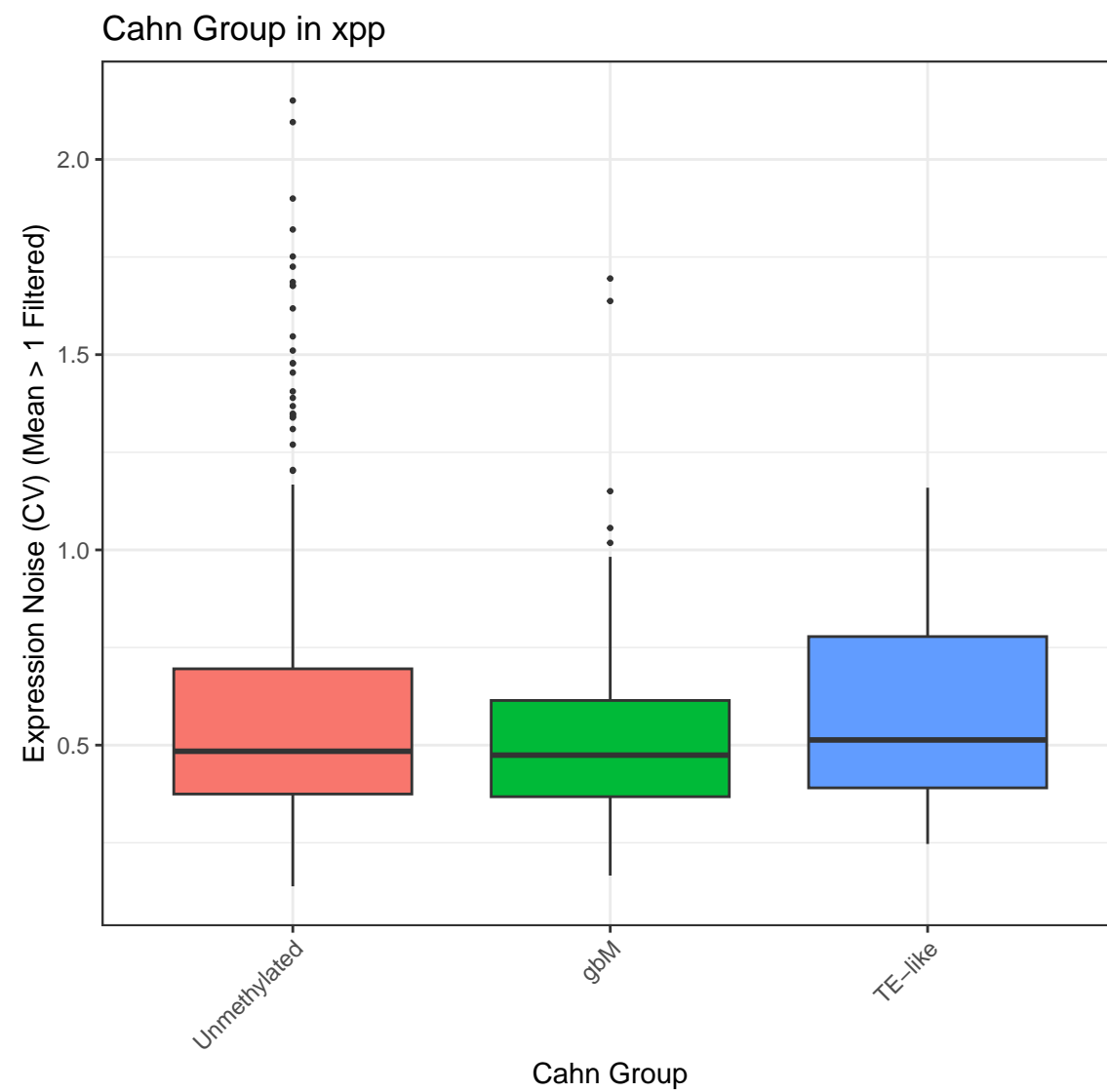
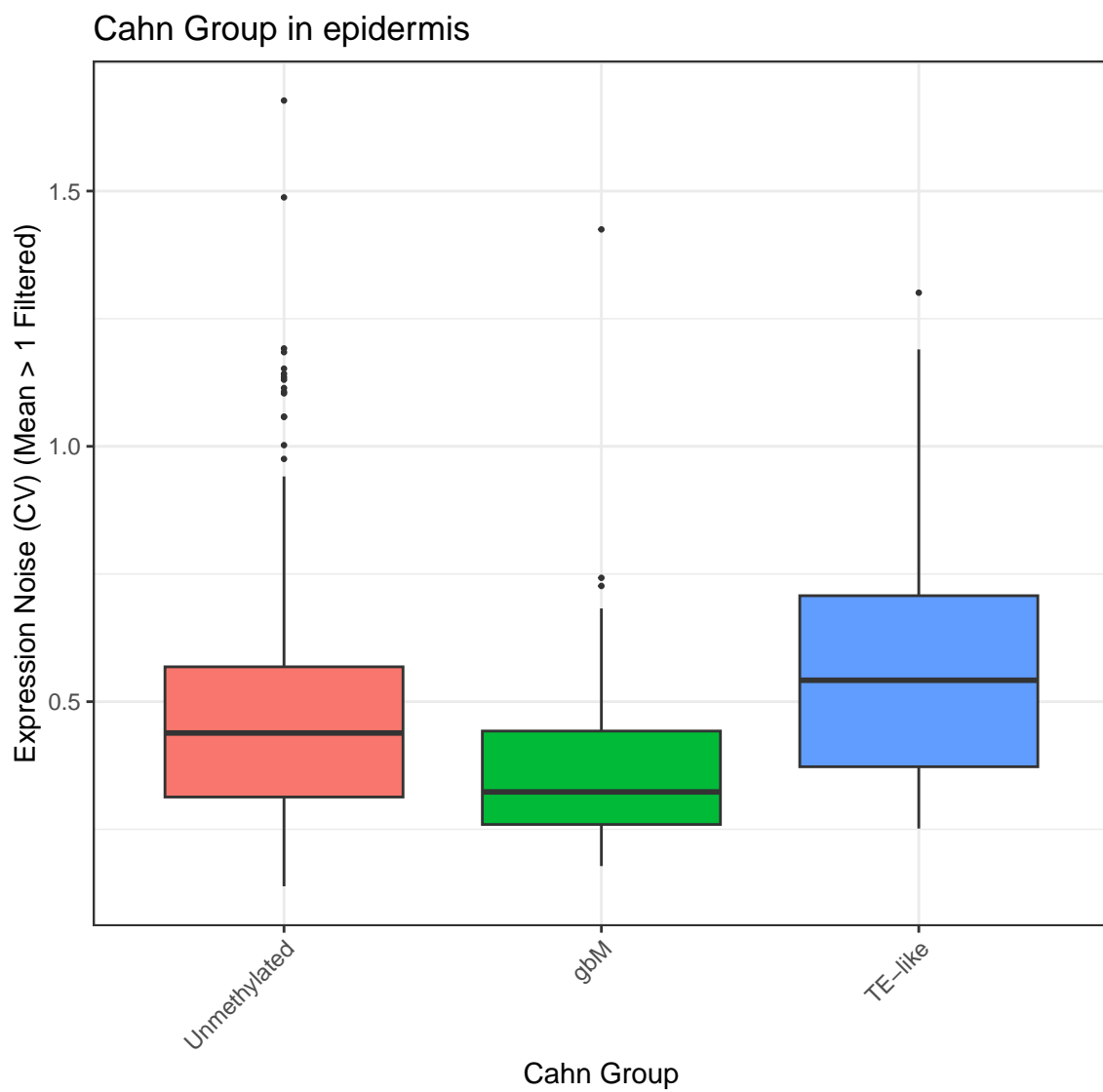
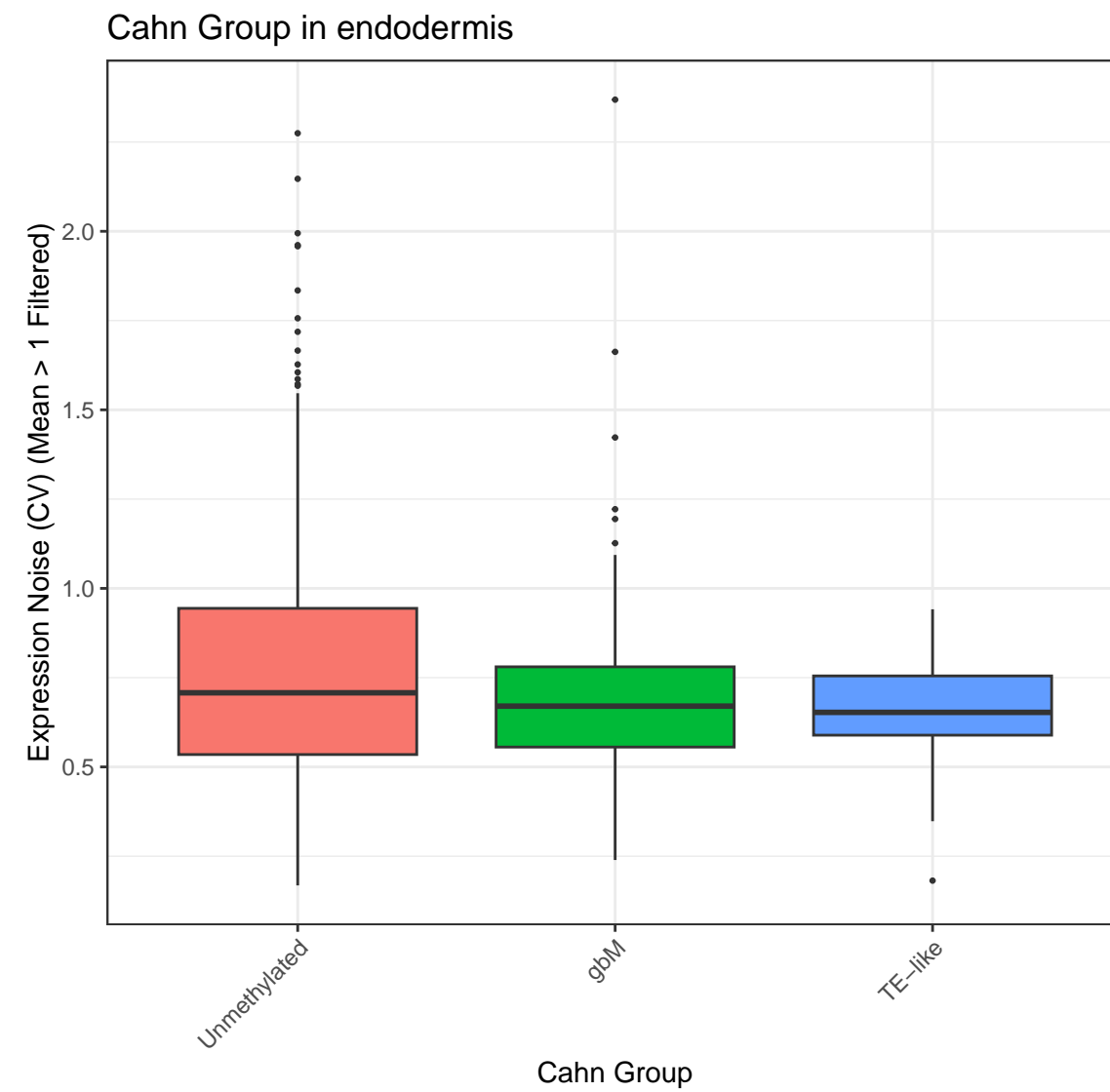
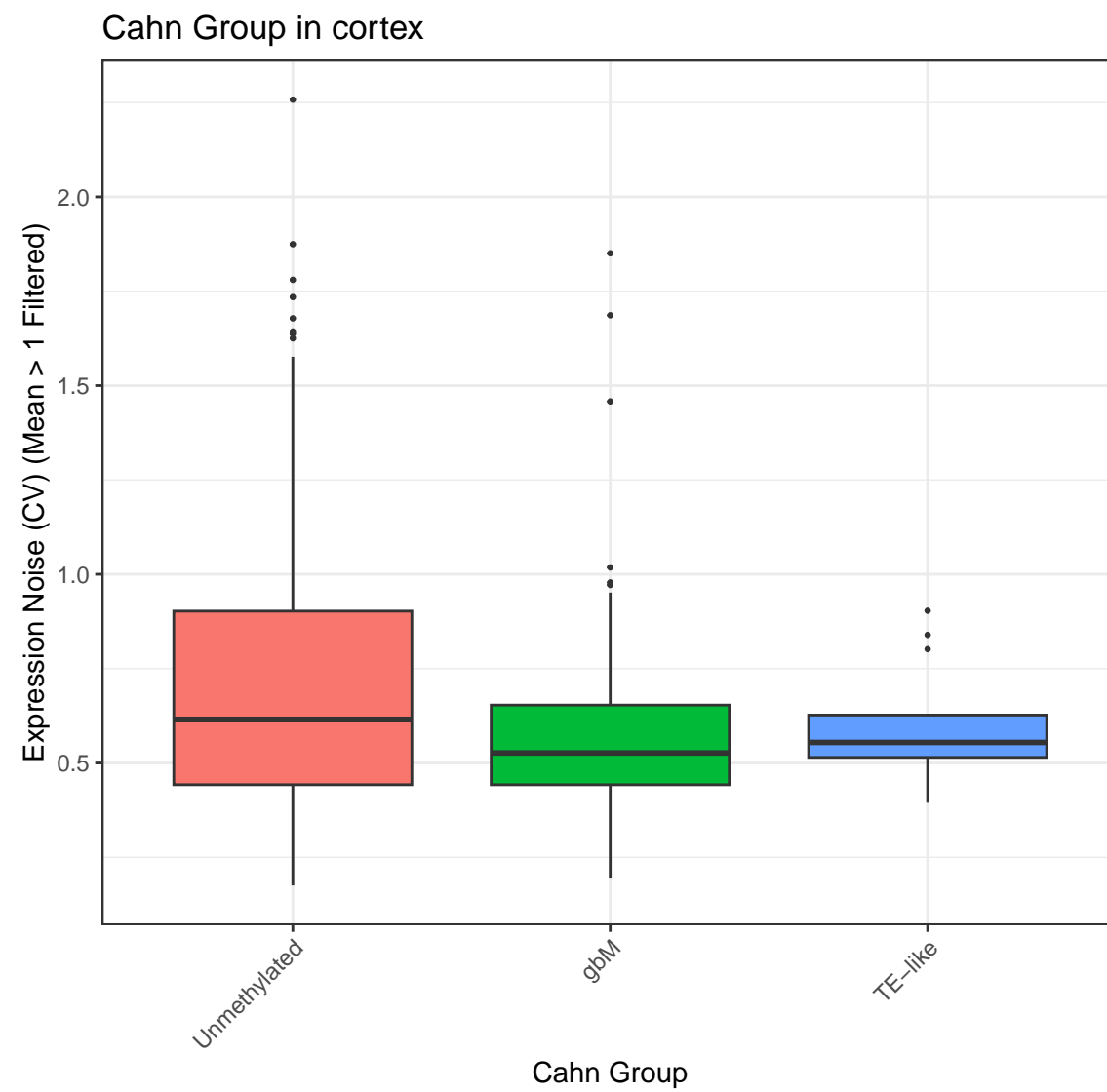
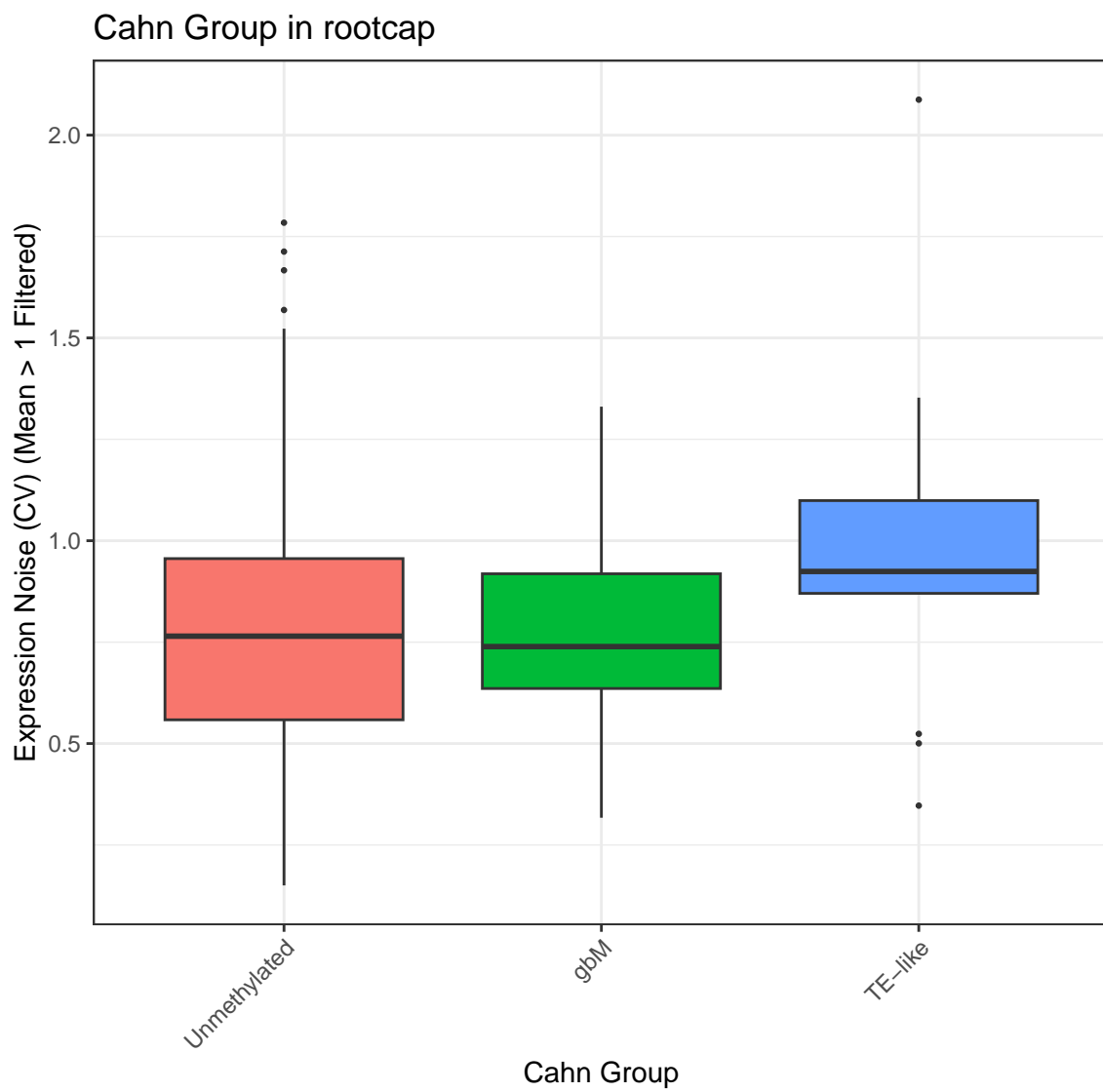
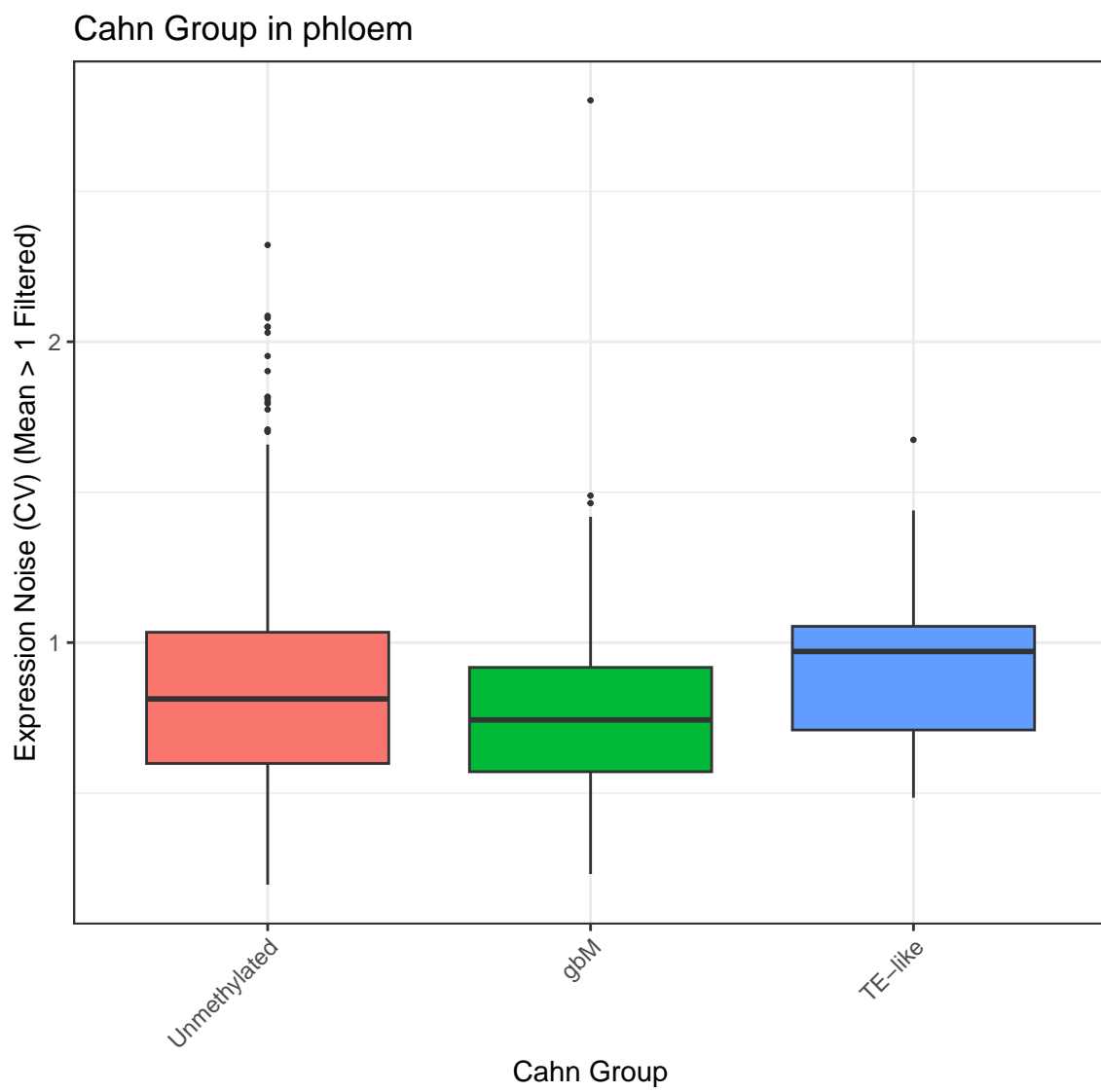
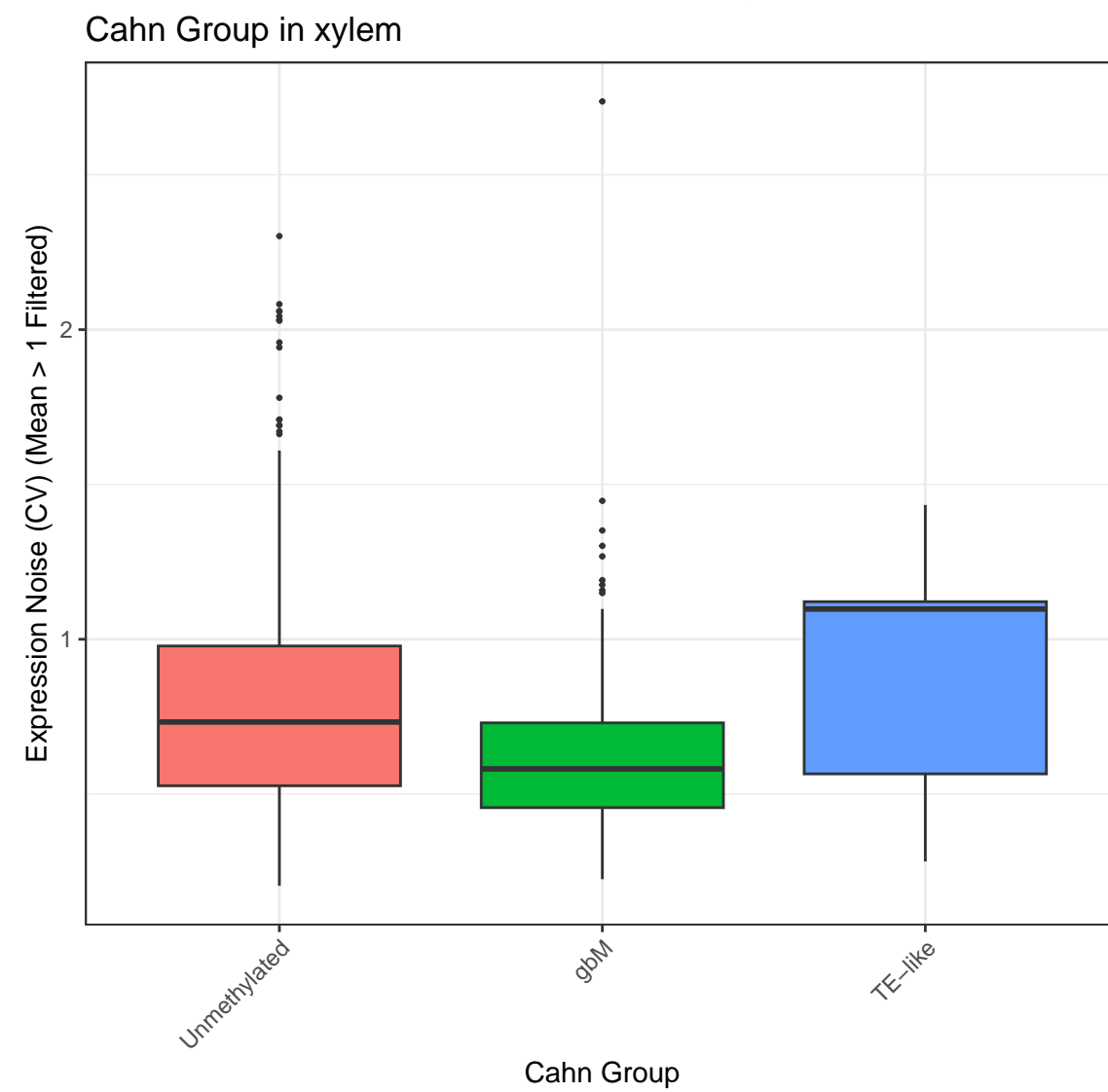
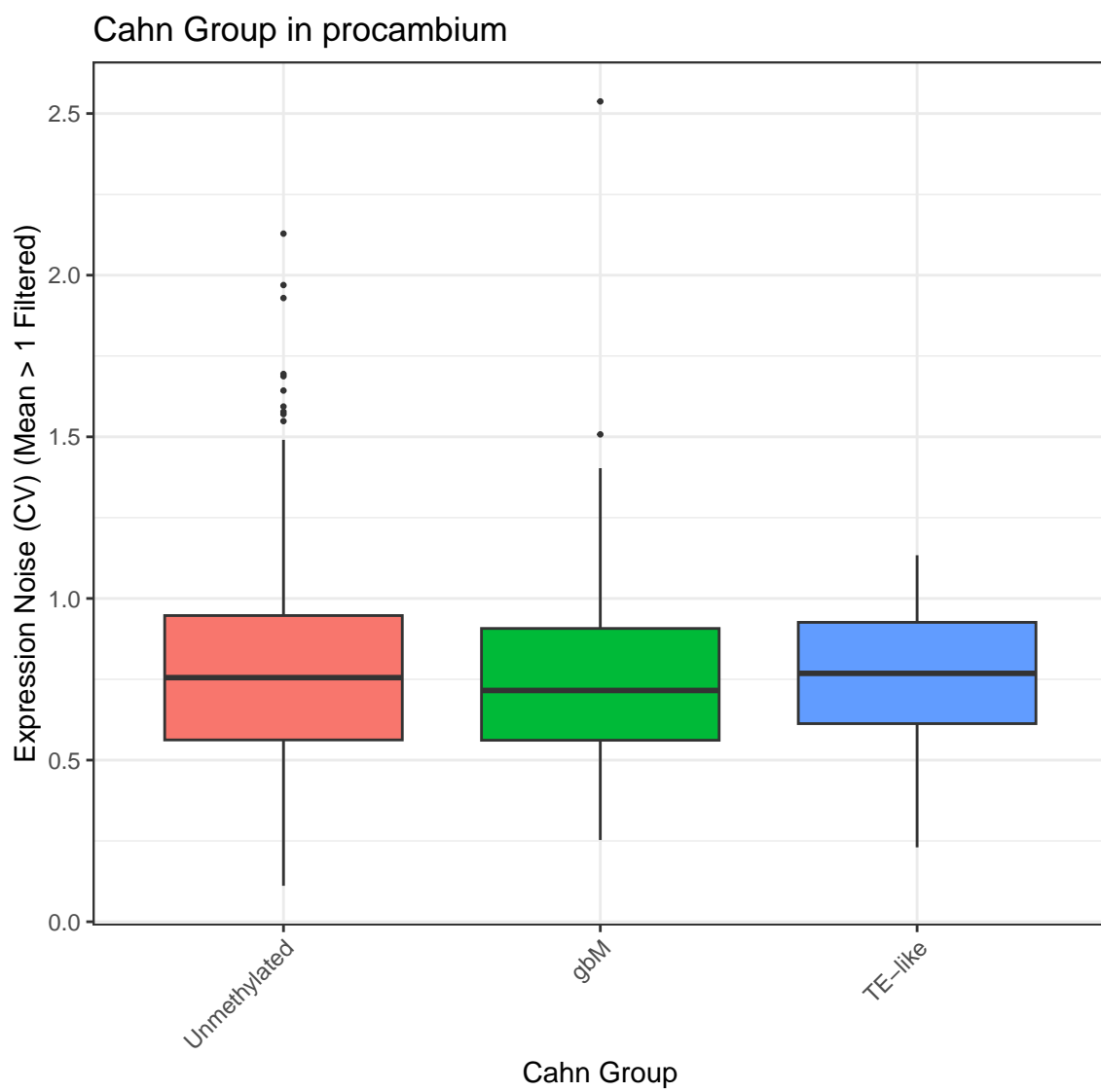




Expression Noise (CV) by Cahn Group in Lineages (Mean &gt; 1 Filtered)



Expression Noise (CV) by Cahn Group in Lineages (Mean > 1 Filtered)



Statistical Comparisons for Expression Noise (CV) by Cahn Group in Lineages (Mean > 1 Filtered)

Context	Filter_Tag	Specific_Context	Group_Type	Comparison	P_value_Formatted
CV Expr by Lineage	Mean > 1 Filtered	rootcap	cahn_group	Overall (Kruskal–Wallis)	9.9792e−02
CV Expr by Lineage	Mean > 1 Filtered	rootcap	cahn_group	gbM vs Unmethylated	5.9868e−01
CV Expr by Lineage	Mean > 1 Filtered	rootcap	cahn_group	TE–like vs Unmethylated	3.8650e−02
CV Expr by Lineage	Mean > 1 Filtered	epidermis	cahn_group	Overall (Kruskal–Wallis)	4.5057e−08
CV Expr by Lineage	Mean > 1 Filtered	epidermis	cahn_group	gbM vs Unmethylated	3.1322e−08
CV Expr by Lineage	Mean > 1 Filtered	epidermis	cahn_group	TE–like vs Unmethylated	1.4365e−01
CV Expr by Lineage	Mean > 1 Filtered	cortex	cahn_group	Overall (Kruskal–Wallis)	6.7060e−04
CV Expr by Lineage	Mean > 1 Filtered	cortex	cahn_group	gbM vs Unmethylated	1.5183e−04
CV Expr by Lineage	Mean > 1 Filtered	cortex	cahn_group	TE–like vs Unmethylated	5.4412e−01
CV Expr by Lineage	Mean > 1 Filtered	xpp	cahn_group	Overall (Kruskal–Wallis)	3.8235e−01
CV Expr by Lineage	Mean > 1 Filtered	xpp	cahn_group	gbM vs Unmethylated	2.3022e−01
CV Expr by Lineage	Mean > 1 Filtered	xpp	cahn_group	TE–like vs Unmethylated	5.7095e−01
CV Expr by Lineage	Mean > 1 Filtered	endodermis	cahn_group	Overall (Kruskal–Wallis)	1.1718e−01
CV Expr by Lineage	Mean > 1 Filtered	endodermis	cahn_group	gbM vs Unmethylated	6.3475e−02
CV Expr by Lineage	Mean > 1 Filtered	endodermis	cahn_group	TE–like vs Unmethylated	2.8405e−01
CV Expr by Lineage	Mean > 1 Filtered	ppp	cahn_group	Overall (Kruskal–Wallis)	9.3234e−03
CV Expr by Lineage	Mean > 1 Filtered	ppp	cahn_group	gbM vs Unmethylated	2.5332e−03
CV Expr by Lineage	Mean > 1 Filtered	ppp	cahn_group	TE–like vs Unmethylated	9.1377e−01
CV Expr by Lineage	Mean > 1 Filtered	procambium	cahn_group	Overall (Kruskal–Wallis)	4.6388e−01
CV Expr by Lineage	Mean > 1 Filtered	procambium	cahn_group	gbM vs Unmethylated	2.5990e−01
CV Expr by Lineage	Mean > 1 Filtered	procambium	cahn_group	TE–like vs Unmethylated	7.0899e−01
CV Expr by Lineage	Mean > 1 Filtered	phloem	cahn_group	Overall (Kruskal–Wallis)	4.3065e−02
CV Expr by Lineage	Mean > 1 Filtered	phloem	cahn_group	gbM vs Unmethylated	6.7639e−02
CV Expr by Lineage	Mean > 1 Filtered	phloem	cahn_group	TE–like vs Unmethylated	1.2611e−01
CV Expr by Lineage	Mean > 1 Filtered	xylem	cahn_group	Overall (Kruskal–Wallis)	3.3365e−06
CV Expr by Lineage	Mean > 1 Filtered	xylem	cahn_group	gbM vs Unmethylated	3.3064e−06
CV Expr by Lineage	Mean > 1 Filtered	xylem	cahn_group	TE–like vs Unmethylated	9.7340e−02