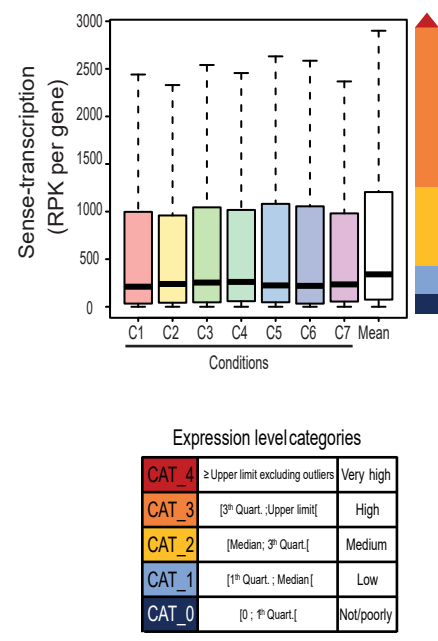
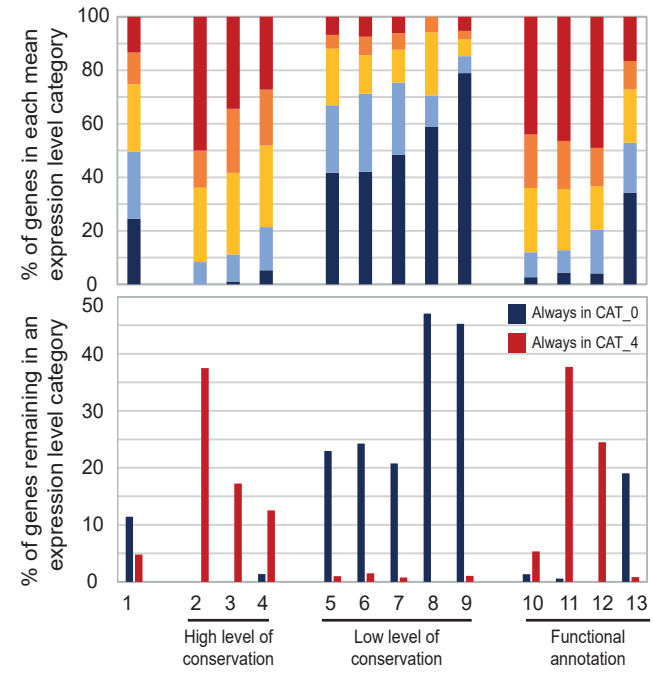


A.

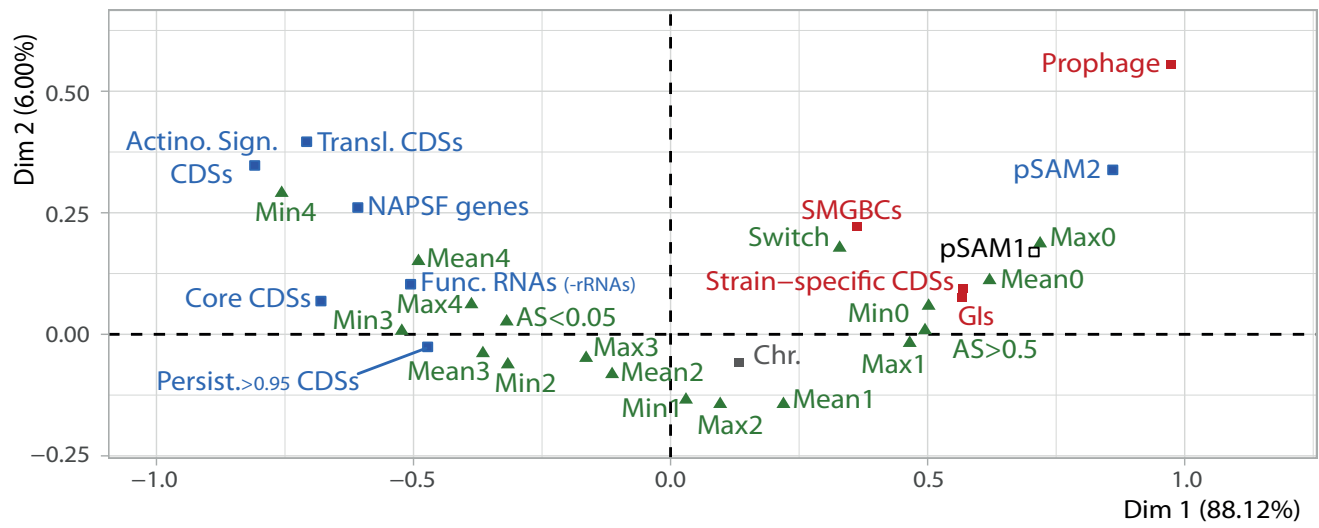


B.

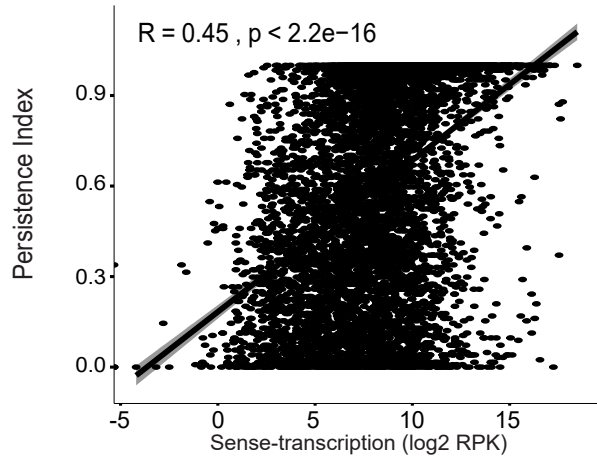


	Proportion of genes always expressed at very high level (CAT_4) compared to the whole set of chromosomal genes		Proportion of genes always not/poorly expressed (CAT_0) compared to the whole set of chromosomal genes	
	Odds ratio	p value	Odds ratio	p value
1. Chr.				
2. Actino.Sig.CDSs	11.9	< 2.2e-16	0.0	2.6e-04
3. Core CDSs	4.2	< 2.2e-16	0.0	< 2.2e-16
4. Persist.>0.95 CDSs	2.9	< 2.2e-16	0.1	< 2.2e-16
5. Gls	0.2	5.5e-12	2.3	< 2.2e-16
6. Strain-specific CDSs	0.3	2.6e-02	2.5	6.1e-07
7. pSAM1	0.2	3.3e-02	2.0	2.2e-03
8. pSAM2	0.0	1	6.9	2.8e-04
9. Prophage	0.2	1.4e-01	6.4	3.0e-16
10. Func. RNAs (-rRNAs)	1.1	7.8e-01	0.1	2.7e-03
11. Transl. CDSs	12.0	< 2.2e-16	0.04	3.2e-08
12. NAPSF genes	6.5	2.9e-06	0.0	5.2e-03
13. SMGBCs	0.2	7.4e-05	1.8	4.7e-05

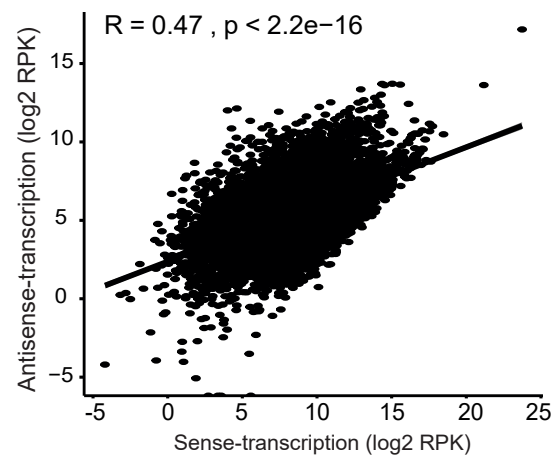
C.



D.



E.



F.

