

4 **S2 Table. General genome features of *S. marcescens* strains 1912768R, 1912768WR and 1912768W.**

Genome feature	Value		
	1912768R	1912768WR	1912768W
Gene number	4927	4998	4997
No. of all scaffolds	1	48	39
Bases in all scaffolds	5117289	5075193	5081421
Gene total length (bp)	4478904	4422783	4450470
Gene average length (bp)	909.05	884.91	890.63
Gene density (kb)	0.96	0.98	0.98
GC content in gene region (%)	61.20	61.25	61.24
Gene/Genome (%)	87.52	87.15	87.58
Intergenic region length (bp)	638385	652410	630951
GC content in intergenic region (%)	49.60	49.88	49.53
Intergenic length/Genome (%)	12.48	12.85	12.42

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