X non-conserved
X similar
X ≥ 70% conserved

	Efm Com15 SagA	Efs 62	Efs OG1RF A	Efs OG1RF B	Efs OG1RF C	Efs D32	Efs DENG1	Efs Symbioflor	Efs 10244	Efs L12	Efs V583	Efs KB1	Efs LD33	Efs L9	Efs Portland ATCC 29212
Efm Com15 SagA	—	42.0	42.0	34.4	56.7	42.0	42.0	42.0	42.0	42.0	42.0	42.0	42.0	42.0	34.4
Efs 62	22.0	—	100.0	36.7	32.6	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	36.7
Efs OG1RF A	22.0	100.0	—	36.7	32.6	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	36.7
Efs OG1RF B	22.9	22.4	22.4	—	31.2	36.7	36.7	36.7	36.7	36.7	36.7	36.7	36.7	36.7	100.0
Efs OG1RF C	37.8	10.2	10.2	10.9	—	32.6	32.6	32.6	32.6	32.6	32.6	32.6	32.6	32.6	31.2
Efs D32	22.0	100.0	100.0	22.4	10.2	—	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	36.7
Efs DENG1	22.0	100.0	100.0	22.4	10.2	100.0	—	100.0	100.0	100.0	100.0	100.0	100.0	100.0	36.7
Efs Symbioflor	22.0	100.0	100.0	22.4	10.2	100.0	100.0	—	100.0	100.0	100.0	100.0	100.0	100.0	36.7
Efs 10244	22.0	100.0	100.0	22.4	10.2	100.0	100.0	100.0	—	100.0	100.0	100.0	100.0	100.0	36.7
Efs L12	22.0	100.0	100.0	22.4	10.2	100.0	100.0	100.0	100.0	—	100.0	100.0	100.0	100.0	36.7
Efs V583	22.0	100.0	100.0	22.4	10.2	100.0	100.0	100.0	100.0	100.0	—	100.0	100.0	100.0	36.7
Efs KB1	22.0	100.0	100.0	22.4	10.2	100.0	100.0	100.0	100.0	100.0	100.0	—	100.0	100.0	36.7
Efs LD33	22.0	100.0	100.0	22.4	10.2	100.0	100.0	100.0	100.0	100.0	100.0	100.0	—	100.0	36.7
Efs L9	22.0	100.0	100.0	22.4	10.2	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	—	36.7
Efs Portland ATCC 29212	22.9	22.4	22.4	100.0	10.9	22.4	22.4	22.4	22.4	22.4	22.4	22.4	22.4	22.4	—

% identity

% similarity