

Interactions	Pull down	Native MS	SEC-MALS	ITC
SpaO/SpaO _C	+	1:2 (2:4)	1:2 (2:4)	ND
SpaO _{V203A} / SpaO _C	+	ND	ND	ND
SpaO _C /SpaO _C	ND	1:1 (2:2, 4:4)	1:1	ND
SpaO ₁₋₁₄₅ /SpaO _C	+	1:2 (1:4)	1:2	K _d = 1.04 ± 0.21 μM
SpaO ₁₋₁₄₅ / SpaO ₁₄₀₋₂₉₇	ND	-	-	ND
SpaO ₁₄₀₋₂₉₇ /SpaO _C	ND	(1:2)	-	ND
SpaO/SpaO _C /OrgB	+	2:4:2 (1:2:2, 1:2:1)	ND	ND
SpaO/SpaO _C /OrgB ₁₋₁₀₅	+	ND	ND	ND
SpaO/SpaO _C /OrgB ₁₀₆₋₂₂₆	+/-	ND	ND	ND
SpaO _{V203A} /OrgB	+	Not stable	ND	ND
OrgB/InvC	+	Not stable	ND	ND
OrgB ₁₋₁₀₅ /InvC	+/-	ND	ND	ND
OrgB ₁₀₆₋₂₂₆ /InvC	+	ND	ND	ND
OrgB/InvC ₁₋₇₉	+	ND	ND	ND
OrgB/InvC ₈₀₋₄₃₁	-	ND	ND	ND
SpaO/SpaO _C /OrgB/InvC	+	1:2:2:1, 2:4:2:1 (2:2:2:1)	2:4:2:1	ND

Table 1. Summary of interacting proteins and domains and complex stoichiometries.

Alternative complexes detected by native mass spectrometry (MS) and SEC-MALS are indicated in brackets. ND= Not determined.