Table S1. Summary of wild-type ParF and mutant ParF<sup>H</sup> protein properties<sup>a</sup>

| Property                           | ParF | ParF <sup>H</sup> |
|------------------------------------|------|-------------------|
| ATPase activity                    | +++  | ++++              |
| ATP binding                        | +++  | ++                |
| ATPase stimulation by ParG         | +++  | -                 |
| Self-association (two hybrid)      | +++  | +++               |
| Interaction with ParG (two hybrid) | +++  | +++               |
| Autopolymerization                 | +++  | +++++             |
| Polymerization inhibition by ADP   | +++  | +++               |
| Polymerization stimulation by ATP  | +++  | -                 |
| Polymerization stimulation by ParG | +++  | +++               |
| Segrosome assembly                 | +++  | +                 |

<sup>&</sup>lt;sup>a</sup>Levels of ParF<sup>H</sup> activity are indicated relative to wild-type activity which is shown as +++