<table>
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<th>method</th>
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<td>overlapping</td>
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<th>comparison</th>
<th>orthologous gene pairs identified by both methods</th>
<th>COG</th>
<th>SOG</th>
<th>% COG</th>
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<tr>
<td>ECO-KPN</td>
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<td>1643</td>
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<td>STM-KPN</td>
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<td>104</td>
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