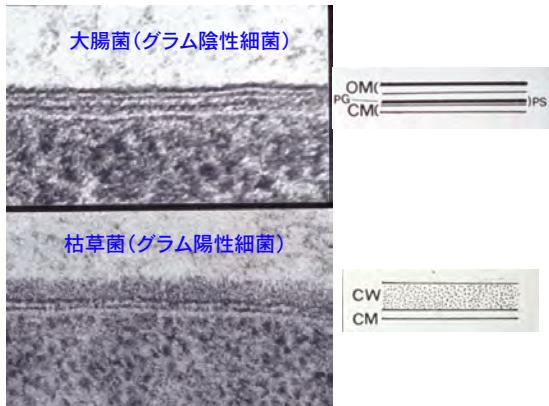
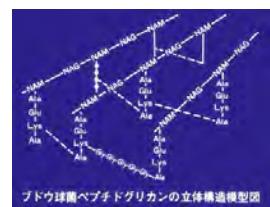
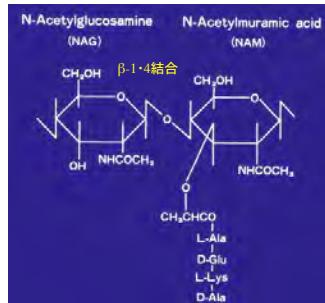


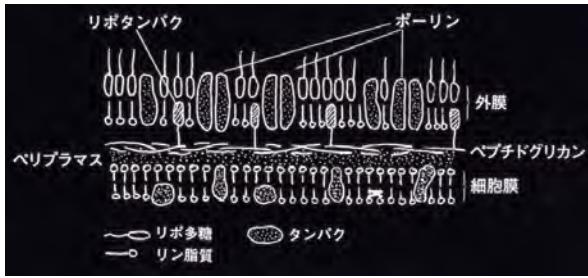
### 7. 超薄切片法で見た細胞壁の構造



### 8. ペプチドグリカンの構成ユニットとペプチド鎖の結合



### 9. グラム陰性外膜

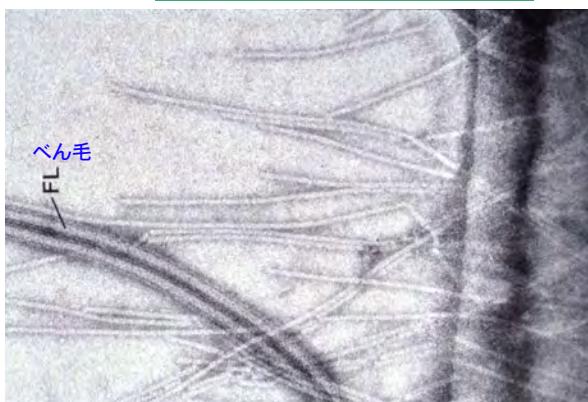


### 15. 毒素原性大腸菌のもつ線毛CFA/I

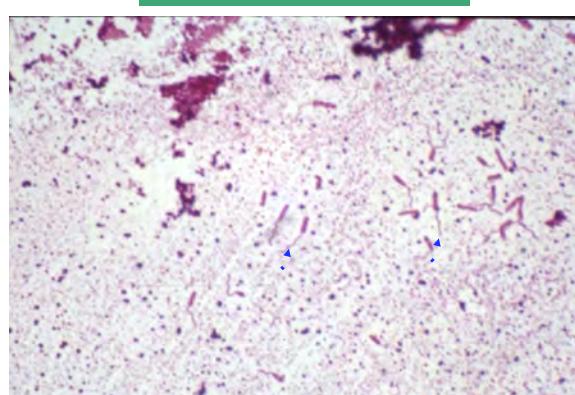
ヒトの腸管粘膜に付着する性質があり腸管への定着因子となっている



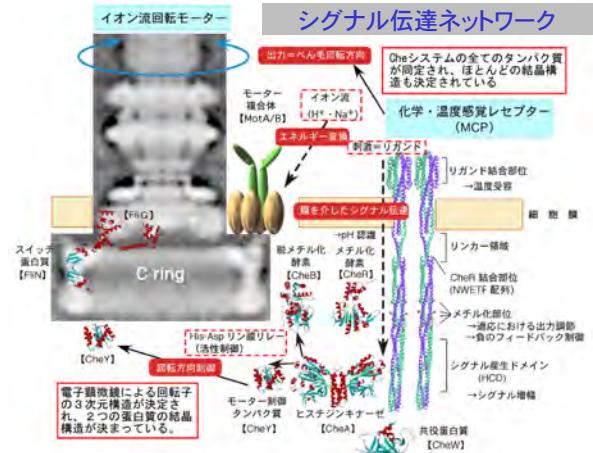
### 16. 大腸菌のtype I 線毛とべん毛



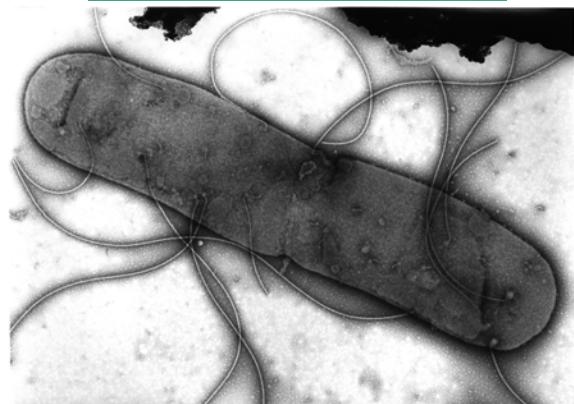
### 17. 緑膿菌のべん毛染色(戸田法)



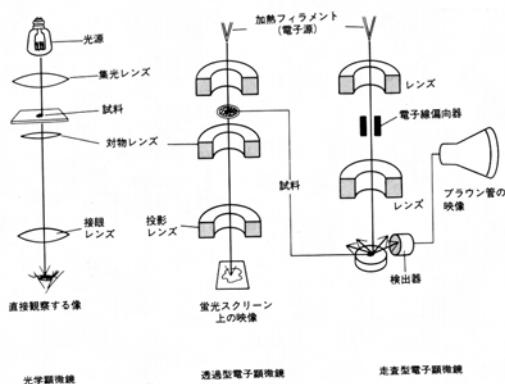
### 18. 菌体へのべん毛のつきかた



Electron micrograph of *Salmonella* cell



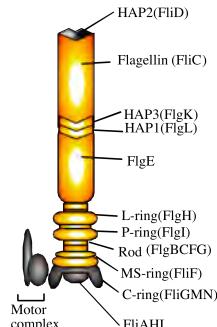
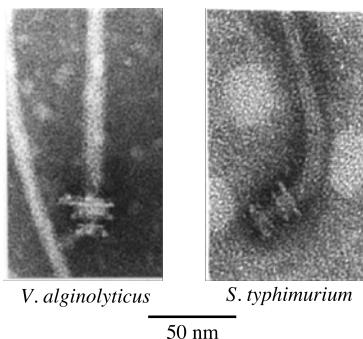
### 電子顕微鏡模式図



最新の電子顕微鏡

安いもので、  
5000万円くらい

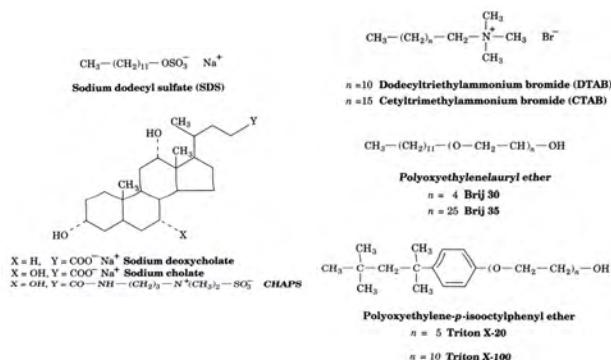
## Flagellar Structure Observed by Electron Microscopy



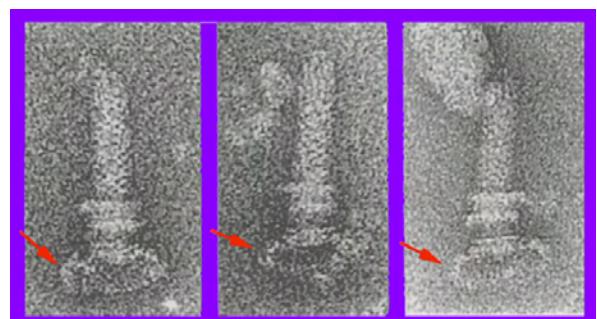
## べん毛の単離プロトコール

- 1) 菌体の回収(低速遠心)
- 2) しょ糖を含む緩衝液にいれる
- 3) リゾチーム処理+ EDTA
- 4) トリトンX-100処理
- 5) MgSO<sub>4</sub>+DNase
- 6) EDTA
- 7) 低速遠心
- 8) 10万gで遠心
- 9) 沈殿をバッファーに懸濁

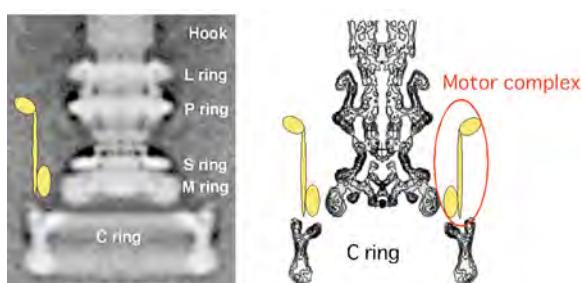
## 生化学で用いられる界面活性剤



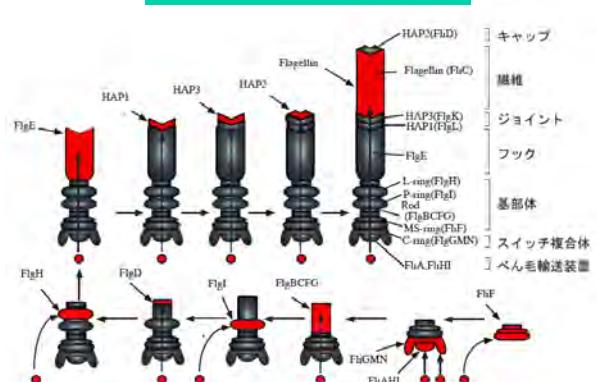
Electron micrographs of hook-basal body and C-ring structure (red arrows)



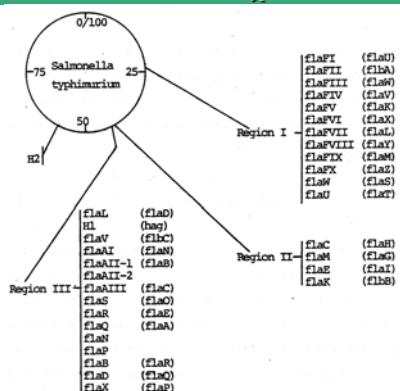
## Motor Structure of flagella



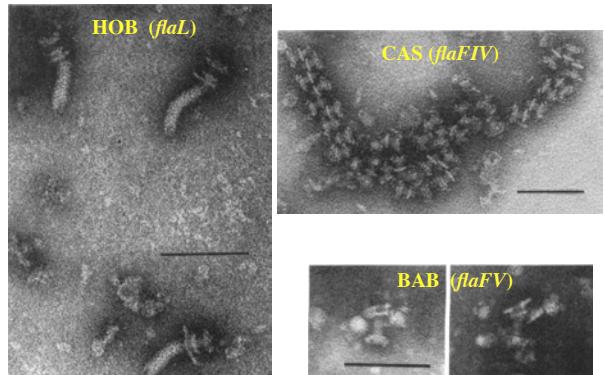
## べん毛形成過程のモデル



**Chromosomal map of the genes essential for flagellar formation in *Salmonella typhimurium*.**

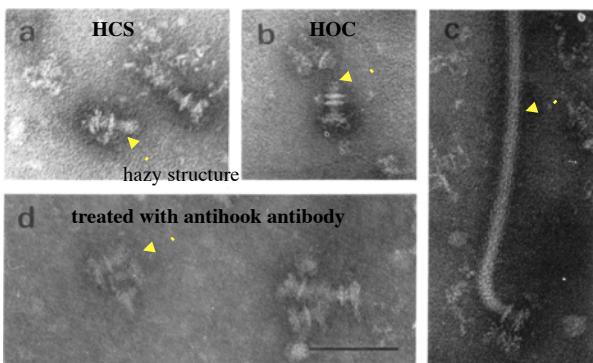


**Flagellar partial structures I**



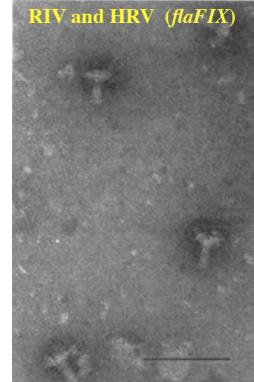
Suzuki et al., J. Bacteriol. (1978)

**Flagellar partial structures II: from a *flaFVIII* mutant**



Suzuki et al., J. Bacteriol. (1978)

**Flagellar partial structures III**



Suzuki et al., J. Bacteriol. (1978)

*Frequencies of flagellar structures detected in nonflagellate mutants<sup>a</sup>*

Determination	Structural entity								
	IF	HOB	BAB	CAS	RIV	RCT	HCS	HRV	HOC
Flagellate parent	++ <sup>b</sup>	+ <sup>c</sup>	+	+ or - <sup>d</sup>	+	+ or -	-	-	-
<i>flaI</i> , <i>flaII</i>	-	++	+	+ or -	+	-	-	-	-
<i>flaL</i>	-	++	+	+ or -	+	-	-	-	-
<i>flaU</i>	-	++	+	+ or -	+	-	-	-	-
<i>flaR</i>	-	++ <sup>e</sup>	+	+ or -	+	-	-	-	-
<i>flaFV</i>	-	-	+	+	++	++	-	-	-
<i>flaFVII</i>	-	-	-	+	-	-	++	+	-
<i>flaF1</i>	-	-	-	-	++	-	-	-	-
<i>flaFIX</i>	-	-	-	-	++	-	-	-	-
<i>flaFIV</i>	-	-	-	-	++	-	-	-	-
<i>flaFVI</i>	-	-	-	-	+	-	-	-	-
<i>flaAI</i>	-	-	-	-	-	-	-	-	-
<i>flaAI</i>	-	-	-	-	-	-	-	-	-
(motC)	++ <sup>f</sup>	+	+	+ or -	+	-	-	-	-
<i>flaAIII</i>	-	-	-	-	-	-	-	-	-
<i>flaB</i>	-	-	-	-	-	-	-	-	-
<i>flaC</i>	-	-	-	-	-	-	-	-	-
<i>flaD</i>	-	-	-	-	-	-	-	-	-
<i>flaE</i>	-	-	-	-	-	-	-	-	-
<i>flaFII</i>	-	-	-	-	-	-	-	-	-
<i>flaFIII</i>	-	-	-	-	-	-	-	-	-
<i>flaFVI</i>	-	-	-	-	-	-	-	-	-
<i>flaFVII</i>	-	-	-	-	-	-	-	-	-
<i>flaFX</i>	-	-	-	-	-	-	-	-	-
<i>flaK</i>	-	-	-	-	-	-	-	-	-
<i>flaM</i>	-	-	-	-	-	-	-	-	-

<sup>a</sup> The frequencies of IF and flagellar basal structures detected in fraction BMII of flagellate parents and nonflagellate mutants are shown.

<sup>b</sup> ++. The count of each structural entity from ½ to 5× the count of IF in its flagellate parent.

<sup>c</sup> +. The count of each structural entity from ½ to ½ the count of IF in its flagellate parent.

<sup>d</sup> -. The count of each structural entity less than ½ the count of IF in its flagellate parent.

<sup>e</sup> Polyhook basal body complexes were detected.

<sup>f</sup> Paralyzed flagella, which were not discriminated morphologically from the IF of their flagellate parent.

**Stepwise process of flagellar morphogenesis in *Salmonella* inferred from the flagellar structures detected on nonflagellate mutants.**

