ERRATUM

Erratum for Hagan et al., "Petrobactin Is Exported from *Bacillus anthracis* by the RND-Type Exporter ApeX"

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Volume 8, no. 5, e01238-17, 2017, https://doi.org/10.1128/mBio.01238-17. Some of the petrobactin component structures in Table 3 were corrupted during file conversion. We apologize for not detecting and correcting this error before publication. The corrected Table 3 is below.

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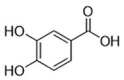
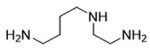
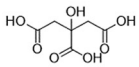
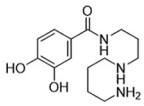
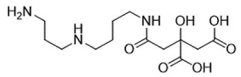
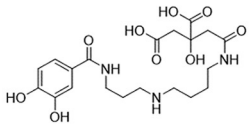
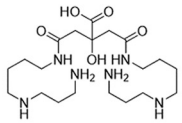
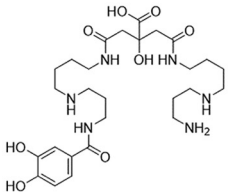
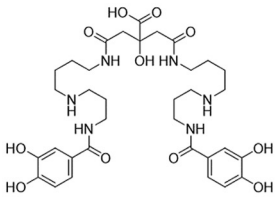
TABLE 3 Structures and molecular weights of predicted petrobactin components present in individual *asbA* to *-F* knockout mutants[a]

Petrobactin Component	Description	Truncated Product (of *asb* mutants)	Size (MolWt)
	3,4-DHB	Δ*asbC*, Δ*asbD*, Δ*asbE* (AsbC-E facilitate 3,4-DHB and spermidine condensation)	154.12
	Spermidine	–	131.22
	Citrate	–	192.12
	3,4-DHB + Spermidine	–	281.36
	Spermidine + Citrate	–	319.36
	3,4-DHB + Spermidine + Citrate	Δ*asbB* (AsbB condenses spermidine to citrate, required for condensation of second spermidine)	455.46
	2(Spermidine) + Citrate	Δ*asbC*, Δ*asbD*, Δ*asbE* (AsbC-E facilitate 3,4-DHB and spermidine condensation) & Δ*asbF* (3,4-DHB synthesis)	446.59
	3,4-DHB + 2(Spermidine) + Citrate	–	582.7
	Petrobactin	Δ*asbA* (AsbB is redundant for AsbA, which condenses citrate to spermidine)	718.81
-	Bacillibactin	–	881.25

[a]Nusca et al. (43).