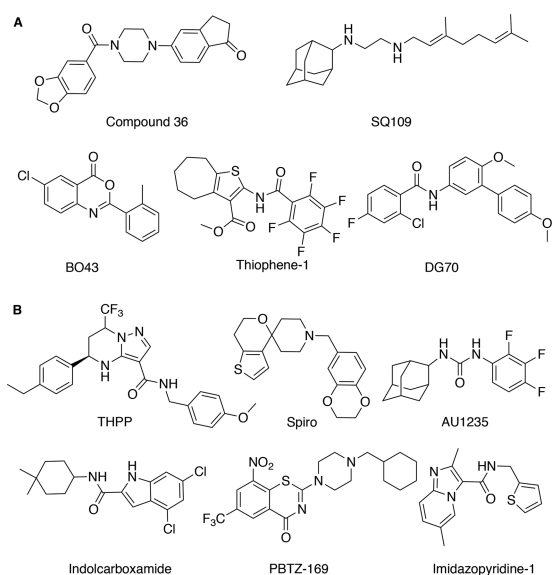


## Correction to Hit Generation in TB Drug Discovery: From Genome to Granuloma

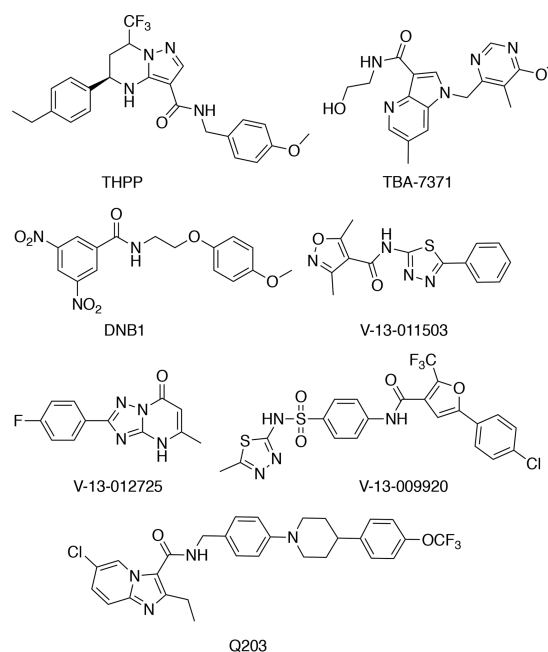
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*Chem. Rev.* 2018, 118 (4), pp 1887–1916. DOI: 10.1021/acs.chemrev.7b00602

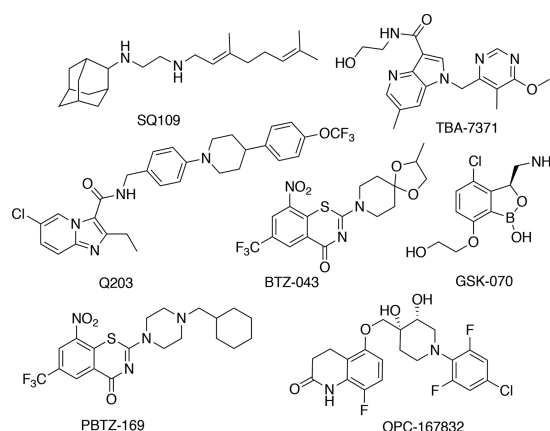
A few structural errors are noticed and listed as follows. A double bond was omitted in the original structure of imidazopyridine-1, TBA-7371, and Q203 in Figures 4, 7, and 8, respectively. The stereochemistry was not illustrated in the original structure of GSK-070 in Figure 8. The full structure of OPC-167832 was not demonstrated in Figure 8. All these errors have been corrected as shown below.



**Figure 4.** Hit compounds (A) identified in target-based whole-cell screens and (B) identified in whole-cell screen under growth culture conditions and discussed in the text. Also see Table 2.



**Figure 7.** Hit compounds identified in intracellular whole-cell screens against *Mtb*-infected macrophages and discussed in the text. Also see Table 2.



**Figure 8.** TB compounds identified in the screens outlined in Table 2 that are currently in clinical trials.