

CORRECTION

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Correction: An asymmetric metal-templated route to amino acids with an isoquinolone core *via* a Rh(III)-catalyzed coupling of aryl hydroxamates with chiral propargylglycine Ni(II) complexes

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Correction for 'An asymmetric metal-templated route to amino acids with an isoquinolone core *via* a Rh(III)-catalyzed coupling of aryl hydroxamates with chiral propargylglycine Ni(II) complexes' by Mikhail A. Arsenov et al., *Org. Biomol. Chem.*, 2022, **20**, 9385–9391, <https://doi.org/10.1039/D2OB01970A>.

The authors regret that there was an error in compound **1g** shown in Scheme 2 (Ar = 3,5-F₂Ph should read Ar = 2,3-F₂Ph). The correct scheme is shown below. The structure has also been corrected in the updated ESI file.

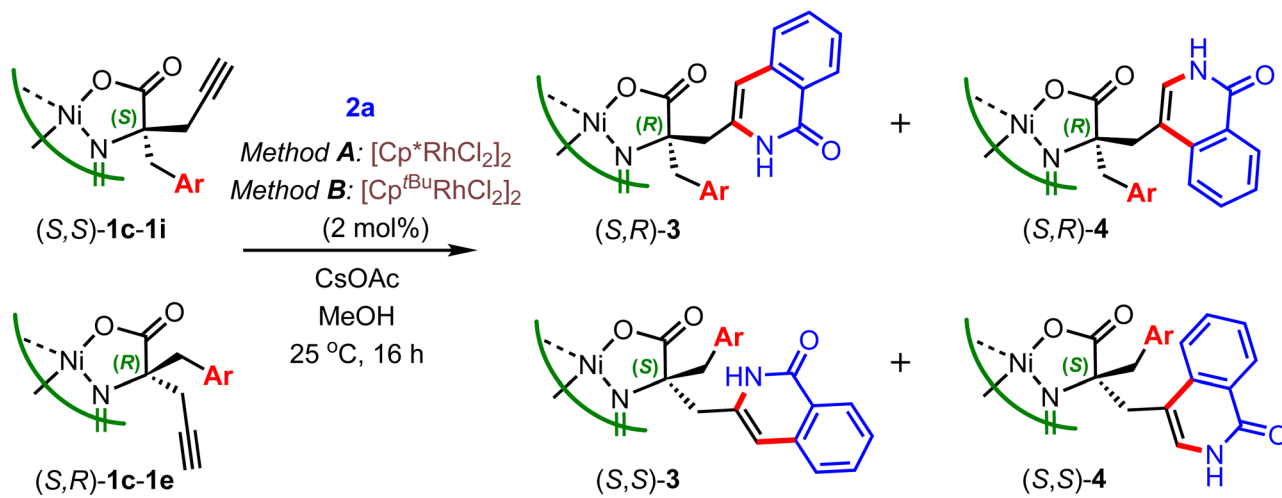
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Ar = Ph (1c):	(S,R)-3ca/(S,R)-4ca 1:1.5 <i>rr</i> (91%) (A); 1:12 <i>rr</i> (93%) (B)
	(S,S)-3ca/(S,S)-4ca 5:1 <i>rr</i> (84%) (A); 1:2 <i>rr</i> (90%) (B)
Ar = 4-<i>t</i>BuPh (1d):	(S,R)-3da/(S,R)-4da 1:1.3 <i>rr</i> (90%) (A); 1:15 <i>rr</i> (92%) (B)
	(S,S)-3da/(S,S)-4da 3.3:1 <i>rr</i> (91%) (A)
Ar = 2-F-4-BrPh (1e):	(S,R)-3ea/(S,R)-4ea 1:2.5 <i>rr</i> (91%) (A)
gram-scale:	0.23 g (22%) + 0.67 g (64%) 1:2.9 <i>rr</i> (A)
	(S,S)-3ea/(S,S)-4ea 3.1:1 <i>rr</i> (87%) (A); 1:2 <i>rr</i> (92%) (B)
Ar = 4-BrPh (1f):	(S,R)-3fa/(S,R)-4fa 1:1.5 <i>rr</i> (90%) (A); 1:13 <i>rr</i> (93%) (B)
Ar = 2,3-F₂Ph (1g):	(S,R)-3ga/(S,R)-4ga 1:2.2 <i>rr</i> (93%) (A); 1:15 <i>rr</i> (95%) (B)
Ar = 4-CF₃Ph (1h):	(S,R)-3ha/(S,R)-4ha 1:1.4 <i>rr</i> (94%) (A); 1:8 <i>rr</i> (93%) (B)
Ar = 4-MeO₂CPh (1i):	(S,R)-3ia/(S,R)-4ia 1:1.5 <i>rr</i> (91%) (A)

Scheme 2 Substrate scope: evaluation of different Phe-derived Ni(II) complexes **1** with phenyl hydroxamate **2a**. Combined isolated yields of the regioisomers **3** and **4** are shown in the parentheses.

The Royal Society of Chemistry apologises for these errors and any consequent inconvenience to authors and readers.

