

Show & Tell



Finally ! This has been a tough nut to crack. I started this project via pairing a Scaleless corn to a Sunglow Motley. It's bad enough that we usually only average about 13 eggs from our mature, veteran het Scaleless females (surely because the eggs are big, from their Emory's Ratsnake ancestry), but factor in the reality that the average number of Scaleless corns is roughly 3.5 per brood, and the odds of getting Scaleless corns with more than one other gene mutation are not yet desirable. Since there are 4 gene mutations in this project (Amel, Motley, Scaleless and Red Factor), the average number of target morphs (in this case, one snake possessing all four gene mutations) is ONE in 32—plus one in 4 on Red Factor?