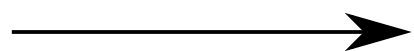


```
{ bionix
, speed ? "normal"
, max-genotypes ? 0
, targets ? [] }:
```



```
{
  options.octopus.call = {
    speed = mkOption {
      type = types.enum [ "normal" "fast" "very-fast" ];
      default = "normal";
    };

    max-genotypes = mkOption {
      type = types.int;
      default = 0;
      description = "Maximum number of candidate genotypes. If
zero then flag is not passed to octopus."
    };

    targets = mkOption {
      type = types.either (types.listOf types.str) types.path;
      default = [];
      description = "either a list of genomic regions to call or a
file containing a list of genomic regions to call";
      example = ''
        targets = [ "chr1" "chr2:0-1,000" ];
      '';
    };
  };
};
...
}
```