

## Product datasheet for **KN201376BN**

### MYH (MUTYH) Human Gene Knockout Kit (CRISPR)

#### Product data:

|               |  |
|---------------|--|
| Product Type: | Knockout Kits (CRISPR)   |
| Format:       | 2 gRNA vectors, 1 mBFP-Neo donor, 1 scramble control   |
| Donor DNA:    | mBFP-Neo   |
| Symbol:       | MYH  |
| Locus ID:     | 4595   |
| Components:   | <b>KN201376G1</b> , MYH gRNA vector 1 in pCas-Guide CRISPR vector (GE100002)<br><b>KN201376G2</b> , MYH gRNA vector 2 in pCas-Guide CRISPR vector (GE100002)<br><b>KN201376BND</b> , donor DNA containing left and right homologous arms and mBFP-Neo functional cassette.<br><b>GE100003</b> , scramble sequence in pCas-Guide vector   |
| Disclaimer:   | These products are manufactured and supplied by OriGene under license from ERS. The kit is designed based on the best knowledge of CRISPR technology. The system has been functionally validated for knocking-in the cassette downstream the native promoter. The efficiency of the knock-out varies due to the nature of the biology and the complexity of the experimental process.  |
| RefSeq:       | <a href="#">NM_001048171</a> , <a href="#">NM_001048172</a> , <a href="#">NM_001048173</a> , <a href="#">NM_001048174</a> , <a href="#">NM_001128425</a> ,<br><a href="#">NM_001293190</a> , <a href="#">NM_001293191</a> , <a href="#">NM_001293192</a> , <a href="#">NM_001293195</a> , <a href="#">NM_001293196</a> ,<br><a href="#">NM_012222</a> , <a href="#">NM_001350650</a> , <a href="#">NM_001350651</a> , <a href="#">NR_146882</a> , <a href="#">NR_146883</a>  |
| UniProt ID:   | <a href="#">Q9UIF7</a>   |
| Synonyms:     | MYH  |
| Summary:      | This gene encodes a DNA glycosylase involved in oxidative DNA damage repair. The enzyme excises adenine bases from the DNA backbone at sites where adenine is inappropriately paired with guanine, cytosine, or 8-oxo-7,8-dihydroguanine, a major oxidatively damaged DNA lesion. The protein is localized to the nucleus and mitochondria. This gene product is thought to play a role in signaling apoptosis by the introduction of single-strand breaks following oxidative damage. Mutations in this gene result in heritable predisposition to colorectal cancer, termed MUTYH-associated polyposis (MAP). Multiple transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Apr 2017] |



[View online »](#)

## Product images:

