

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 9, 2026

## hMSH2 Clone E3 (full-length)

RRID:Addgene\_16453

Type: Plasmid

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### Proper Citation

RRID:Addgene\_16453

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### Plasmid Information

**URL:** <http://www.addgene.org/16453>

**Proper Citation:** RRID:Addgene\_16453

**Insert Name:** MSH2

**Organism:** Homo sapiens

**Bacterial Resistance:** Ampicillin

**Defining Citation:** [PMID:8261515](#)

**Vector Backbone Description:** Backbone Marker:Stratagene; Backbone Size:3000; Vector Backbone:pBluescript SK; Vector Types:Bacterial Expression; Bacterial Resistance:Ampicillin

**Comments:** A full-length clone of hMSH2 cDNA was cloned into the XbaI/XhoI site of pBluescript SK.

**Plasmid Name:** hMSH2 Clone E3 (full-length)

**Record Creation Time:** 20220422T221929+0000

**Record Last Update:** 20260808T081133+0000

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### Ratings and Alerts

No rating or validation information has been found for hMSH2 Clone E3 (full-length).

No alerts have been found for hMSH2 Clone E3 (full-length).

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### Data and Source Information

**Source:** [Addgene](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [PRECISE-TBI](#) .

Ryu SW, et al. (2020) Proteome-wide identification of HSP70/HSC70 chaperone clients in human cells. PLoS biology, 18(7), e3000606.