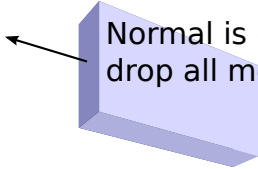


Normal is closest to z-axis:
drop all monomials with z factors.



Normal is closest to (minus) x-axis:
drop all monomials with x factors.