

Ingredient
A : int Ci : int Hf Hi : int Ni : int Oi : int allIngr : dict elements flag : str lastLine : int lineIngr : dict lineNo : NoneType, int name rho
find(name) getLine(lineNo) nitrocellulose(nitration) readFile(fileName)

Mixture
Ci : int Delta : float Hf : int Hi : int Ni : int Oi : int Tv b compoDict f gamma its : int n name p rho speciesList tol_b : float tol_z : float
balanceAt(T, verbose, its, tol) prettyPrint()

Reaction
LHS RHS name
Hr(R) Kp(T)

Specie
Hf__R : NoneType allFits : dict cardNo date fitThigh fitFlow fitTmid formula molWt : NoneType, int phase specieName
Cp(T, R) Cv(T, R) G(T, R) H(T, R, C) S(T, R, C) dumpJSON() get(specieName) getMMH(T, R, C, ref) read(curvePath)