

# Generalized Linear Models

Generalized linear models currently supports estimation using the one-parameter exponential families.

See [Module Reference](#) [#module-reference] for commands and arguments.

## Examples

```
# Load modules and data
In [1]: import statsmodels.api as sm

In [2]: data = sm.datasets.scotland.load(as_pandas=False)

In [3]: data.exog = sm.add_constant(data.exog)

# Instantiate a gamma family model with the default link function.
In [4]: gamma_model = sm.GLM(data.endog, data.exog, family=sm.families.Gamma())

In [5]: gamma_results = gamma_model.fit()

In [6]: print(gamma_results.summary())
```

Generalized Linear Model Regression Results

Dep. Variable:	y	No. Observations:	32
Model:	GLM	Df Residuals:	24
Model Family:	Gamma	Df Model:	7
Link Function:	inverse_power	Scale:	0.0035843
Method:	IRLS	Log-Likelihood:	-83.017
Date:	Tue, 02 Feb 2021	Deviance:	0.087389
Time:	07:07:06	Pearson chi2:	0.0860
No. Iterations:	6		
Covariance Type:	nonrobust		

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	coef	std err	z	P> z	[0.025	0.975]
const	-0.0178	0.011	-1.548	0.122	-0.040	0.005
x1	4.962e-05	1.62e-05	3.060	0.002	1.78e-05	8.14e-05
x2	0.0020	0.001	3.824	0.000	0.001	0.003
x3	-7.181e-05	2.71e-05	-2.648	0.008	-0.000	-1.87e-05
x4	0.0001	4.06e-05	2.757	0.006	3.23e-05	0.000
x5	-1.468e-07	1.24e-07	-1.187	0.235	-3.89e-07	9.56e-08
x6	-0.0005	0.000	-2.159	0.031	-0.001	-4.78e-05
x7	-2.427e-06	7.46e-07	-3.253	0.001	-3.89e-06	-9.65e-07

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Detailed examples can be found here:

- [GLM](#) [examples/notebooks/generated/glm.html]
- [Formula](#) [examples/notebooks/generated/glm\_formula.html]

## Technical Documentation

The statistical model for each observation  $i$  is assumed to be

$$Y_i \sim F_{EDM}(\cdot|\theta, \phi, w_i) \text{ and } \mu_i = E[Y_i|x_i] = g^{-1}(x_i'\beta).$$

where  $g$  is the link function and  $F_{EDM}(\cdot|\theta, \phi, w)$  is a distribution of the family of exponential dispersion models (EDM) with natural parameter  $\theta$ , scale parameter  $\phi$  and weight  $w$ . Its density is given by

$$f_{EDM}(y|\theta, \phi, w) = c(y, \phi, w) \exp\left(\frac{y\theta - b(\theta)}{\phi} w\right).$$

It follows that  $\mu = b'(\theta)$  and  $Var[Y|x] = \frac{\phi}{w} b''(\theta)$ . The inverse of the first equation gives the natural parameter as a function of the expected value  $\theta(\mu)$  such that

$$Var[Y_i|x_i] = \frac{\phi}{w_i} v(\mu_i)$$

with  $v(\mu) = b''(\theta(\mu))$ . Therefore it is said that a GLM is determined by link function  $g$  and variance function  $v(\mu)$  alone (and  $x$  of course).

Note that while  $\phi$  is the same for every observation  $y_i$  and therefore does not influence the estimation of  $\beta$ , the weights  $w_i$  might be different for every  $y_i$  such that the estimation of  $\beta$  depends on them.

Distribution	Domain	$\mu = E[Y x]$	$v(\mu)$	$\theta(\mu)$	$b(\theta)$	$\phi$
Binomial $B(n, p)$	$0, 1, \dots, n$	$np$	$\mu - \frac{\mu^2}{n}$	$\log \frac{p}{1-p}$	$n \log(1 + e^\theta)$	1
Poisson $P(\mu)$	$0, 1, \dots, \infty$	$\mu$	$\mu$	$\log(\mu)$	$e^\theta$	1
Neg. Binom. $NB(\mu, \alpha)$	$0, 1, \dots, \infty$	$\mu$	$\mu + \alpha\mu^2$	$\log(\frac{\alpha\mu}{1+\alpha\mu})$	$-\frac{1}{\alpha} \log(1 - \alpha e^\theta)$	1
Gaussian/Normal $N(\mu, \sigma^2)$	$(-\infty, \infty)$	$\mu$	1	$\mu$	$\frac{1}{2}\theta^2$	$\sigma^2$
Gamma $N(\mu, \nu)$	$(0, \infty)$	$\mu$	$\mu^2$	$-\frac{1}{\mu}$	$-\log(-\theta)$	$\frac{1}{\nu}$
Inv. Gauss. $IG(\mu, \sigma^2)$	$(0, \infty)$	$\mu$	$\mu^3$	$-\frac{1}{2\mu^2}$	$-\sqrt{-2\theta}$	$\sigma^2$
Tweedie $p \geq 1$	depends on $p$	$\mu$	$\mu^p$	$\frac{\mu^{1-p}}{1-p}$	$\frac{\alpha-1}{\alpha} \left(\frac{\theta}{\alpha-1}\right)^\alpha$	$\phi$

The Tweedie distribution has special cases for  $p = 0, 1, 2$  not listed in the table and uses  $\alpha = \frac{p-2}{p-1}$ .

Correspondence of mathematical variables to code:

- $Y$  and  $y$  are coded as `endog`, the variable one wants to model
- $x$  is coded as `exog`, the covariates alias explanatory variables
- $\beta$  is coded as `params`, the parameters one wants to estimate
- $\mu$  is coded as `mu`, the expectation (conditional on  $x$ ) of  $Y$
- $g$  is coded as `link` argument to the `class Family`
- $\phi$  is coded as `scale`, the dispersion parameter of the EDM
- $w$  is not yet supported (i.e.  $w = 1$ ), in the future it might be `var_weights`
- $p$  is coded as `var_power` for the power of the variance function  $v(\mu)$  of the Tweedie distribution, see table
- $\alpha$  is either
  - Negative Binomial: the ancillary parameter `alpha`, see table
  - Tweedie: an abbreviation for  $\frac{p-2}{p-1}$  of the power  $p$  of the variance function, see table

## References

- Gill, Jeff. 2000. Generalized Linear Models: A Unified Approach. SAGE QASS Series.
- Green, P.J. 1984. "Iteratively reweighted least squares for maximum likelihood estimation, and some robust and resistant alternatives." Journal of the Royal Statistical Society, Series B, 46, 149-192.

- Hardin, J.W. and Hilbe, J.M. 2007. "Generalized Linear Models and Extensions." 2nd ed. Stata Press, College Station, TX.
- McCullagh, P. and Nelder, J.A. 1989. "Generalized Linear Models." 2nd ed. Chapman & Hall, Boca Rotan.

## Module Reference

### Model Class

[GLM](#)

[generated/statsmodels.genmod.generalized\_linear\_model.GLM.html#statsmodels.genmod.generalized\_linear\_model.GLM](e

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### Results Class

[GLMResults](#) [generated/statsmodels.genmod.generalized\_linear\_model.GLMResults.html#statsmodels.genmod.generalized\_li

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[PredictionResults](#)

[generated/statsmodels.genmod.generalized\_linear\_model.PredictionResults.html#statsmodels.genmod.generalized\_linear\_n

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### Families

The distribution families currently implemented are

[Family](#) [generated/statsmodels.genmod.families.family.Family.html#statsmodels.genmod.families.family.Family](link, varianc

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[Binomial](#) [generated/statsmodels.genmod.families.family.Binomial.html#statsmodels.genmod.families.family.Binomial]([link)

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[Gamma](#) [generated/statsmodels.genmod.families.family.Gamma.html#statsmodels.genmod.families.family.Gamma]([link])

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[Gaussian](#) [generated/statsmodels.genmod.families.family.Gaussian.html#statsmodels.genmod.families.family.Gaussian]([link)

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[InverseGaussian](#)

[generated/statsmodels.genmod.families.family.InverseGaussian.html#statsmodels.genmod.families.family.InverseGaussian

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[NegativeBinomial](#)

[generated/statsmodels.genmod.families.family.NegativeBinomial.html#statsmodels.genmod.families.family.NegativeBinomi

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[Poisson](#) [generated/statsmodels.genmod.families.family.Poisson.html#statsmodels.genmod.families.family.Poisson]([link])

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[Tweedie](#) [generated/statsmodels.genmod.families.family.Tweedie.html#statsmodels.genmod.families.family.Tweedie]([link, va

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## Link Functions

The link functions currently implemented are the following. Not all link functions are available for each distribution family. The list of available link functions can be obtained by

```
>>> sm.families.family.<familyname>.links
```

[Link](#) [generated/statsmodels.genmod.families.links.Link.html#statsmodels.genmod.families.links.Link]()

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[CDFLink](#) [generated/statsmodels.genmod.families.links.CDFLink.html#statsmodels.genmod.families.links.CDFLink]([dbn])

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[CLogLog](#) [generated/statsmodels.genmod.families.links.CLogLog.html#statsmodels.genmod.families.links.CLogLog]()

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[Log](#) [generated/statsmodels.genmod.families.links.Log.html#statsmodels.genmod.families.links.Log]()

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[Logit](#) [generated/statsmodels.genmod.families.links.Logit.html#statsmodels.genmod.families.links.Logit]()

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[NegativeBinomial](#)

[generated/statsmodels.genmod.families.links.NegativeBinomial.html#statsmodels.genmod.families.links.NegativeBinomial]

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[Power](#) [generated/statsmodels.genmod.families.links.Power.html#statsmodels.genmod.families.links.Power]([power])

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[cauchy](#) [generated/statsmodels.genmod.families.links.cauchy.html#statsmodels.genmod.families.links.cauchy]()

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[cloglog](#) [generated/statsmodels.genmod.families.links.cloglog.html#statsmodels.genmod.families.links.cloglog]()

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[identity](#) [generated/statsmodels.genmod.families.links.identity.html#statsmodels.genmod.families.links.identity]()

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[inverse\\_power](#)

[generated/statsmodels.genmod.families.links.inverse\_power.html#statsmodels.genmod.families.links.inverse\_power]()

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[inverse\\_squared](#)

[generated/statsmodels.genmod.families.links.inverse\_squared.html#statsmodels.genmod.families.links.inverse\_squared]()

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[log](#) [generated/statsmodels.genmod.families.links.log.html#statsmodels.genmod.families.links.log]()

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[logit](#) [generated/statsmodels.genmod.families.links.logit.html#statsmodels.genmod.families.links.logit]()

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[nbinom](#) [generated/statsmodels.genmod.families.links.nbinom.html#statsmodels.genmod.families.links.nbinom]([alpha])

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[probit](#) [generated/statsmodels.genmod.families.links.probit.html#statsmodels.genmod.families.links.probit]([dbn])

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## Variance Functions

Each of the families has an associated variance function. You can access the variance functions here:

```
>>> sm.families.<familyname>.variance
```

[VarianceFunction](#)

[generated/statsmodels.genmod.families.varfuncs.VarianceFunction.html#statsmodels.genmod.families.varfuncs.VarianceFunction]

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[constant](#) [generated/statsmodels.genmod.families.varfuncs.constant.html#statsmodels.genmod.families.varfuncs.constant]

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[Power](#) [generated/statsmodels.genmod.families.varfuncs.Power.html#statsmodels.genmod.families.varfuncs.Power]([power],

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[mu](#) [generated/statsmodels.genmod.families.varfuncs.mu.html#statsmodels.genmod.families.varfuncs.mu]

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[mu\\_squared](#) [generated/statsmodels.genmod.families.varfuncs.mu\_squared.html#statsmodels.genmod.families.varfuncs.mu\_squared]

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[mu\\_cubed](#) [generated/statsmodels.genmod.families.varfuncs.mu\_cubed.html#statsmodels.genmod.families.varfuncs.mu\_cubed]

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[Binomial](#) [generated/statsmodels.genmod.families.varfuncs.Binomial.html#statsmodels.genmod.families.varfuncs.Binomial]

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[binary](#) [generated/statsmodels.genmod.families.varfuncs.binary.html#statsmodels.genmod.families.varfuncs.binary]

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[NegativeBinomial](#)

[generated/statsmodels.genmod.families.varfuncs.NegativeBinomial.html#statsmodels.genmod.families.varfuncs.NegativeBinomial]

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[nbinom](#) [generated/statsmodels.genmod.families.varfuncs.nbinom.html#statsmodels.genmod.families.varfuncs.nbinom]

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