

MODELING DARK ENERGY WITH GENETIC ALGORITHMS

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Outline/Summary

In collaboration with Savvas Nesseris [1205.0364]

- A decade since the discovery of acceleration, and we still don't know what Dark Energy is!**
- We need a model-independent and bias-free reconstruction of the observables that we deduce from data.**
- We propose a new method: Genetic Algorithms**
- Error estimation with Path Integral formalism**
- Show how it works for S_nIa, BAO & growth data**
- Conclusions: Λ CDM works pretty well...**

How can we infer the DE dynamics?

- Characterize **observables** as **functions** of redshift
- Operate on functions (**differentiate, integrate**)
- Search for **relationships** between functions
- Derive other **physical quantities** from them
- Look for **consistency relations**
- Only then **compare** with Dark Energy models

Genetic Algorithm does that for you!

Evolution at work!

How does evolution work?

- Individuals have a **genome** (a pool of genes)
- The genome mutates and mixes (via sex)
- There is a rate of **mutation** and **crossover**
- Next generation's genome is different
- **Natural selection** “crops the tree”: **speciation**
- Only a few individuals (species) survive
- Some species live for a long time (**survival**)

How do Genetic Algorithms work?

Population
(individuals)



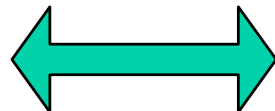
Functions
(cont. & differen.)

Generations
(genome)



Grammar
(poly, exp, trig, log)

Fitness (survival)



χ^2 (minimized)

Mutations (rand.)



Operations (rand.)

Crossover (sex)



Recombination

Lineage (species)



Convergence (limit)

How do Genetic Algorithms work?

- The grammar and the allowed operations determines how the Genetic Algorithm works
- There are two important parameters:
 - The **mutation rate** (5 – 10%) is the probability that an arbitrary part of a genome changes
 - The **selection rate** (10 – 20 %) is the fraction of individuals that finally produce offsprings
- If rates are **larger** then **GA may not converge**
- If rates are **smaller** then **GA converges too slowly**
- Important: jumps between different minima, in order to explore the whole space of functions

Advantages of Genetic Algorithms

- No “a priori” assumptions on the DE model (e.g. # free parameters)
- In general DE models are not well understood (parameter space too large or too complex)
- **OUR GOAL: minimize a functional $\chi^2(f(z))$ thru stochastic processes based on GA evolution.**
- Similar to a **Path Integral**, but where dynamics is dictated by the data w/o ‘a priori’ assumptions.
- Use the **DATA** to find the **MODEL**
- The end result doesn’t have any free parameters

An example of Genetic Algorithm

- Assume our **grammar** includes **basic functions**:
polynomials, trigonometric, exponentials, logs, ...

1.- Set up random initial population $M(0)$, e.g.

$$f_{GA,1}(x) = \ln x$$

$$f_{GA,2}(x) = -1 + x + x^2$$

$$f_{GA,3}(x) = \sin x$$

[Note: The number of functions in a generic population is typically of a **few hundreds!**]

An example of Genetic Algorithm

2.- Given some data points w/ errors (x_i, y_i, σ_i) , the algorithm measures the **fitness** of each function

$$\chi^2(f) \equiv \sum_{i=1}^N \left(\frac{y_i - f(x_i)}{\sigma_i} \right)^2$$

In our example:

$$\chi_1^2(x) = 200$$

$$\chi_2^2(x) = 500$$

$$\chi_3^2(x) = 1000$$

An example of Genetic Algorithm

3.- Selection is done via **“tournament method”**, sorting population w.r.t. fitness χ^2 each individual

A fixed % total population, adjusted by selection rate, are chosen out (in our example only 2 of 3).

Reproduction = crossover & mutation

- **Crossover** will randomly combine parts of parent solutions

$$\begin{aligned} f_{GA,1}(x) \oplus f_{GA,2}(x) &\rightarrow (\bar{f}_{GA,1}(x), \bar{f}_{GA,2}(x), \bar{f}_{GA,3}(x)) \\ &= (\ln(x^2), -1 + \ln(x^2), -1 + \ln(x)) \end{aligned}$$

An example of Genetic Algorithm

- **Mutation** will randomly change the functions, according to mutation rate, e.g. exponent of some terms or the value of a coefficient.

$$\bar{f}_{GA,3}(x) = -1 + \ln(x) \rightarrow -1 + \ln(x^3)$$

Then the next generation becomes:

$$M(1) = (\bar{f}_{GA,1}(x), \bar{f}_{GA,2}(x), \bar{f}_{GA,3}(x)) = (\ln(x^2), -1 + \ln(x^2), -1 + \ln(x^3))$$

An example of Genetic Algorithm

3.- Back to beginning: evaluate fitness χ^2 of each individual:

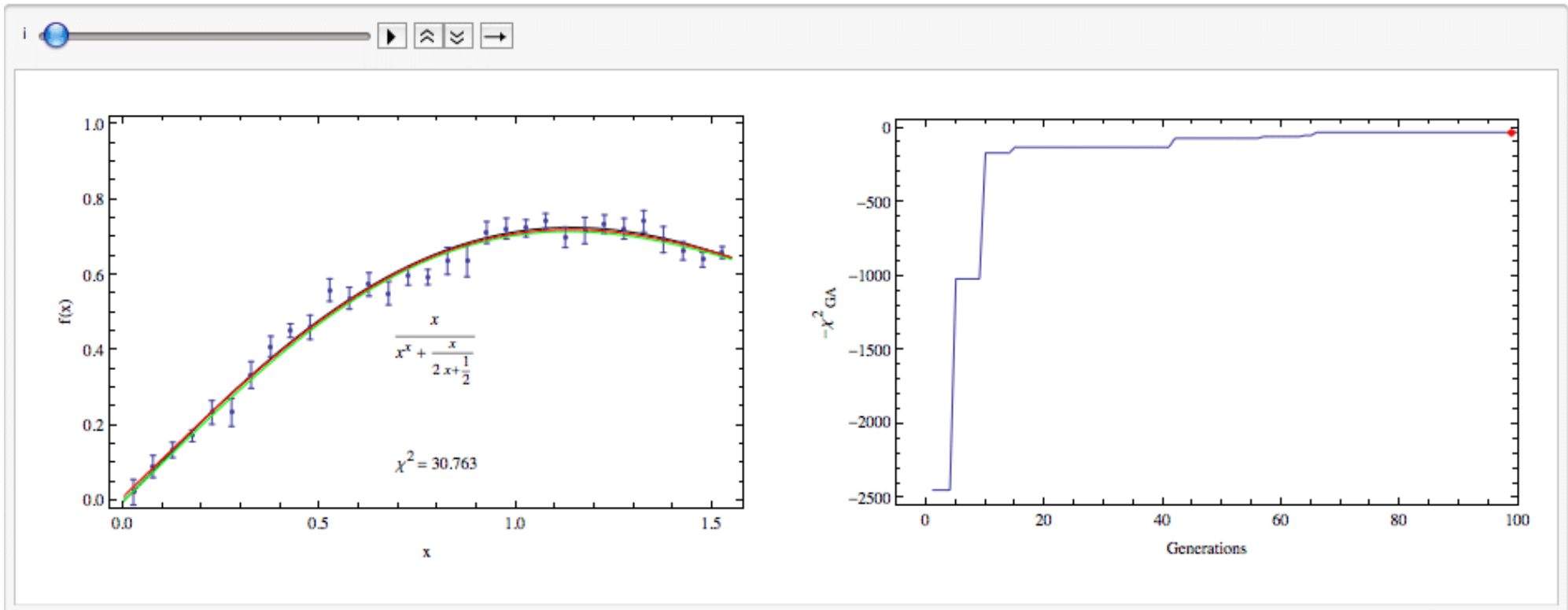
$$\chi_1^2(x) = 150, \quad \chi_2^2(x) = 300, \quad \chi_3^2(x) = 400$$

Even after a single generation, the combined χ^2 of each individual has decreased significantly!

It usually takes \sim 100 generations to reach a reasonable χ^2 and \sim 500 generations to converge to a minimum χ^2 , and \sim 1000 generations until a predetermined goal has been reached.

An example of Genetic Algorithm

Input function $f(x) = \frac{1}{4} + \left(x - \frac{1}{4}\right) \exp\left[-\frac{1}{2}x^2\right]$



Best χ^2 function $f_{best \chi^2}(x) = x / \left(x^x + 2x / (4x + 1)\right)$

Error estimation for GAs

Construct a functional $\mathcal{L} = \mathcal{N} \exp(-\chi^2(f)/2)$

where $\chi^2(f) \equiv \sum_{i=1}^N \left(\frac{y_i - f(x_i)}{\sigma_i} \right)^2$

Properly normalized functional

$$\int \mathcal{D}f \mathcal{L} = \int \mathcal{D}f \mathcal{N} \exp(-\chi^2(f)/2) = 1$$

where $\mathcal{N} = \left((2\pi)^{N/2} \prod_{i=1}^N \sigma_i \right)^{-1}$

Error estimation for GAs

The 1σ error around the best-fit function (BFF)

$$\begin{aligned} CI(x_i, \delta f_i) &= \int_{f_{bf}(x_i) - \delta f_i}^{f_{bf}(x_i) + \delta f_i} df_i \frac{1}{(2\pi)^{1/2} \sigma_i} \exp\left(-\frac{1}{2} \left(\frac{y_i - f_i}{\sigma_i}\right)^2\right) \\ &= \frac{1}{2} \left(\operatorname{erf}\left(\frac{\delta f_i + f_{bf}(x_i) - y_i}{\sqrt{2}\sigma_i}\right) + \operatorname{erf}\left(\frac{\delta f_i - f_{bf}(x_i) + y_i}{\sqrt{2}\sigma_i}\right) \right) \\ &= \operatorname{erf}\left(1/\sqrt{2}\right) \quad \text{at each point } \mathbf{x}_i \end{aligned}$$

Then minimize to get $\delta f(\mathbf{x})$ – a band around BFF

$$\chi_{CI}^2(\delta f_i) = \sum_{i=1}^N \left(CI(x_i, \delta f_i) - \operatorname{erf}\left(1/\sqrt{2}\right) \right)^2$$

Concrete Example for DE

Data sets and models

Dataset	Information	FRW/DE models	LTB models
SnIa	$d_L(z)$	$(1+z) \int_0^z \frac{dx}{H(x)/H_0}$	$(1+z)^2 A(r(z), t(z))$
BAO	$H(z), d_A(z)$	$\frac{l_{BAO}(z \text{ drag})}{D_V(z)}$	$(1+z)\xi(z) \frac{l_{BAO}(r_\infty, t_0)}{D_V(z)}$ ^a
Growth-rate $f = \frac{d \ln(\delta)}{d \ln a}$	$\gamma, \Omega_m(a)$	$\Omega_m(a) = \frac{\Omega_{0m} a^{-3}}{H(a)^2/H_0^2}$	$(1 - \Omega_M^{-1}(r)) \frac{A(r, t)}{r}$ ^b

^awhere in both cases $D_V(z) = \left((1+z)^2 D_A(z)^2 \frac{z}{H(z)} \right)^{1/3}$ and $H(z) = H_L(z)$ for the LTB

^bwhere $\Omega_M(r)$ is the mass radial function.

Genetic Algorithm data functions

SnIa – luminosity distance

$$GA_1(z) \equiv D_L(z) = d_L(z)/H_0^{-1}$$

BAO – angular diameter distance

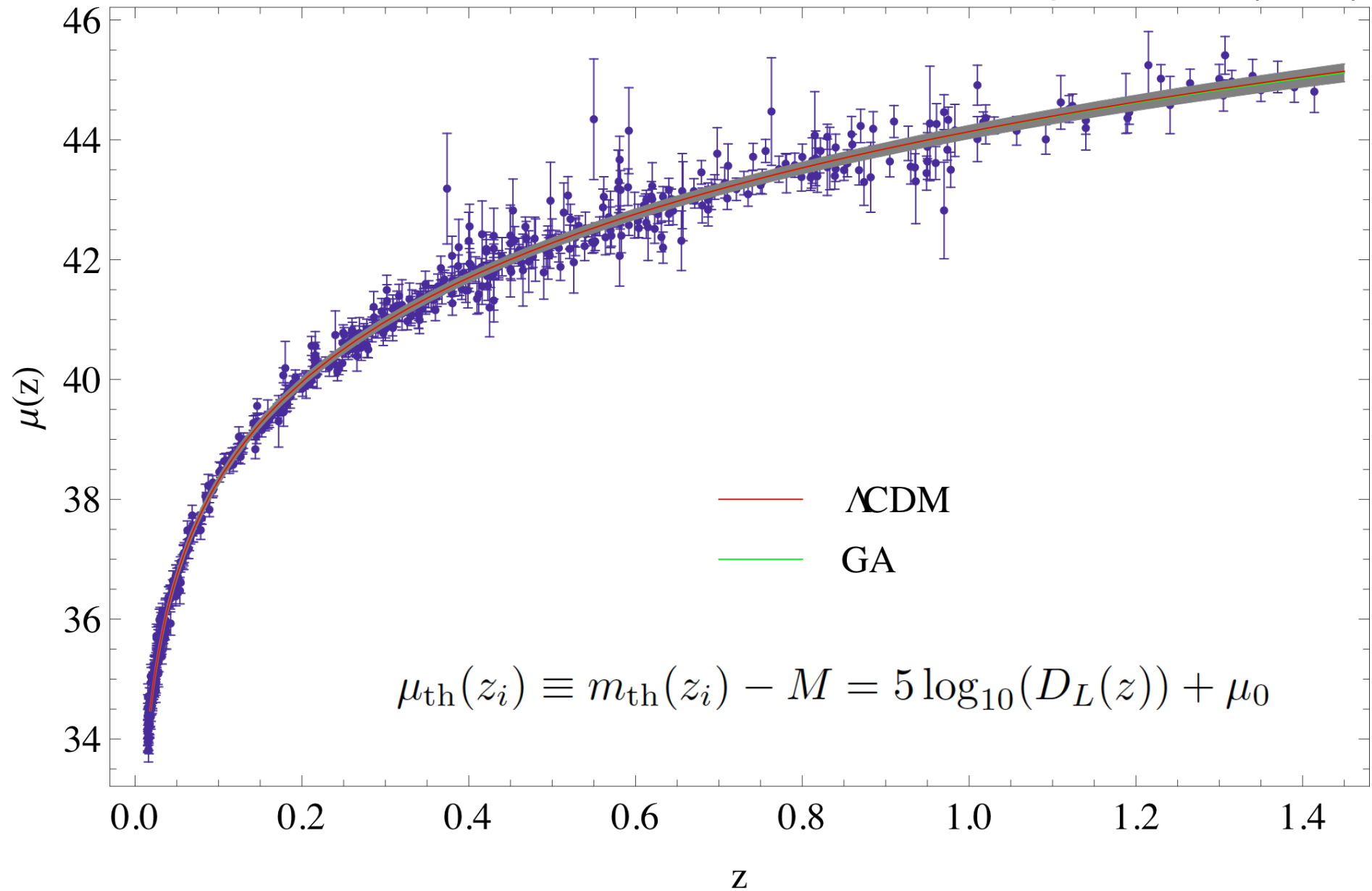
$$d_z(z) = \frac{l_{BAO}(z_{drag})}{D_V(z)} = \frac{l_{BAO}(\Omega_{0m} = 1)}{z} GA_2(z)$$

Growth rate of matter structures

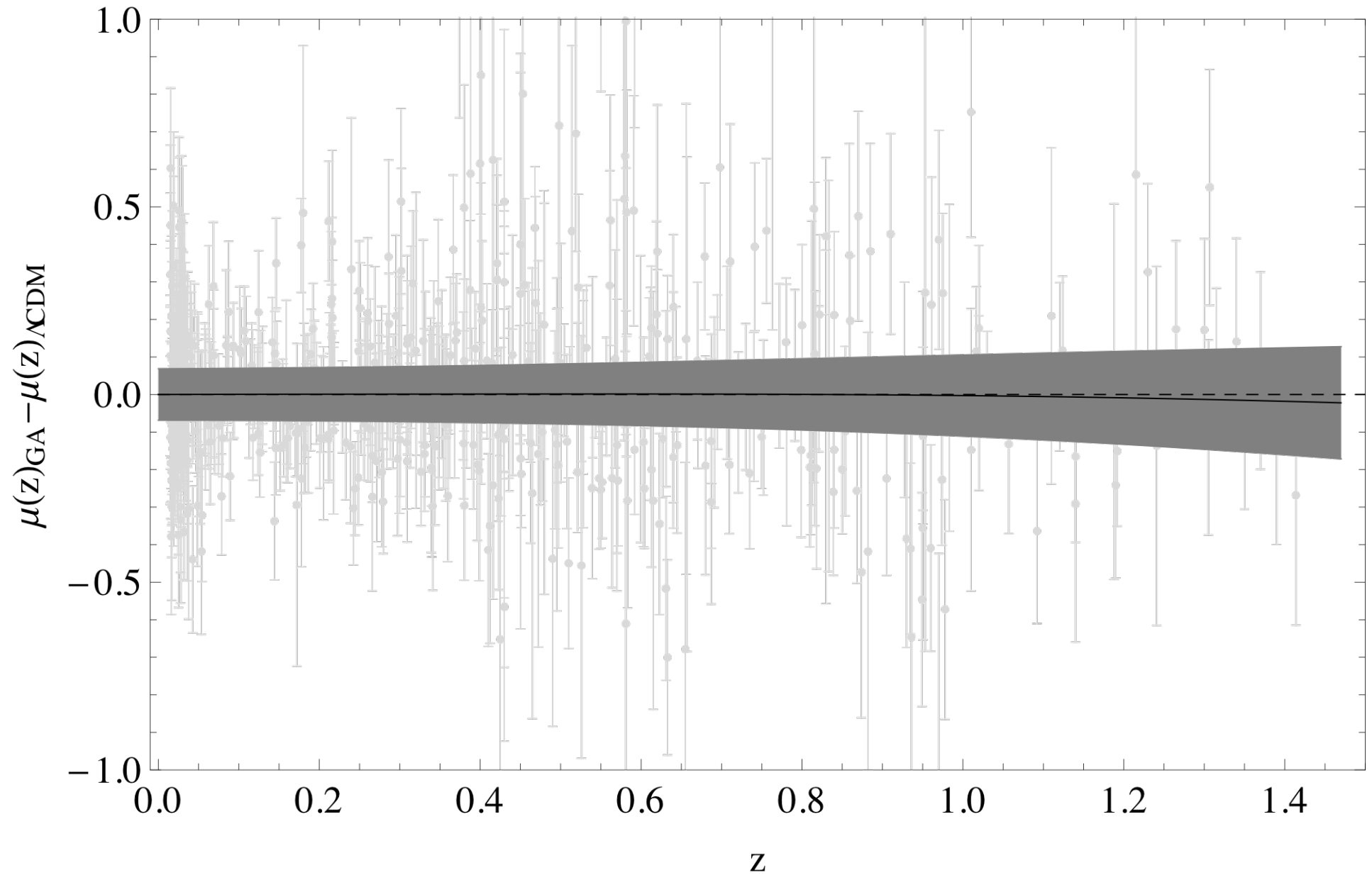
$$f\sigma_8(z) \equiv f(z) \cdot \sigma_8(z) = GA_3(z)$$

Genetic Algorithms & SNIa data

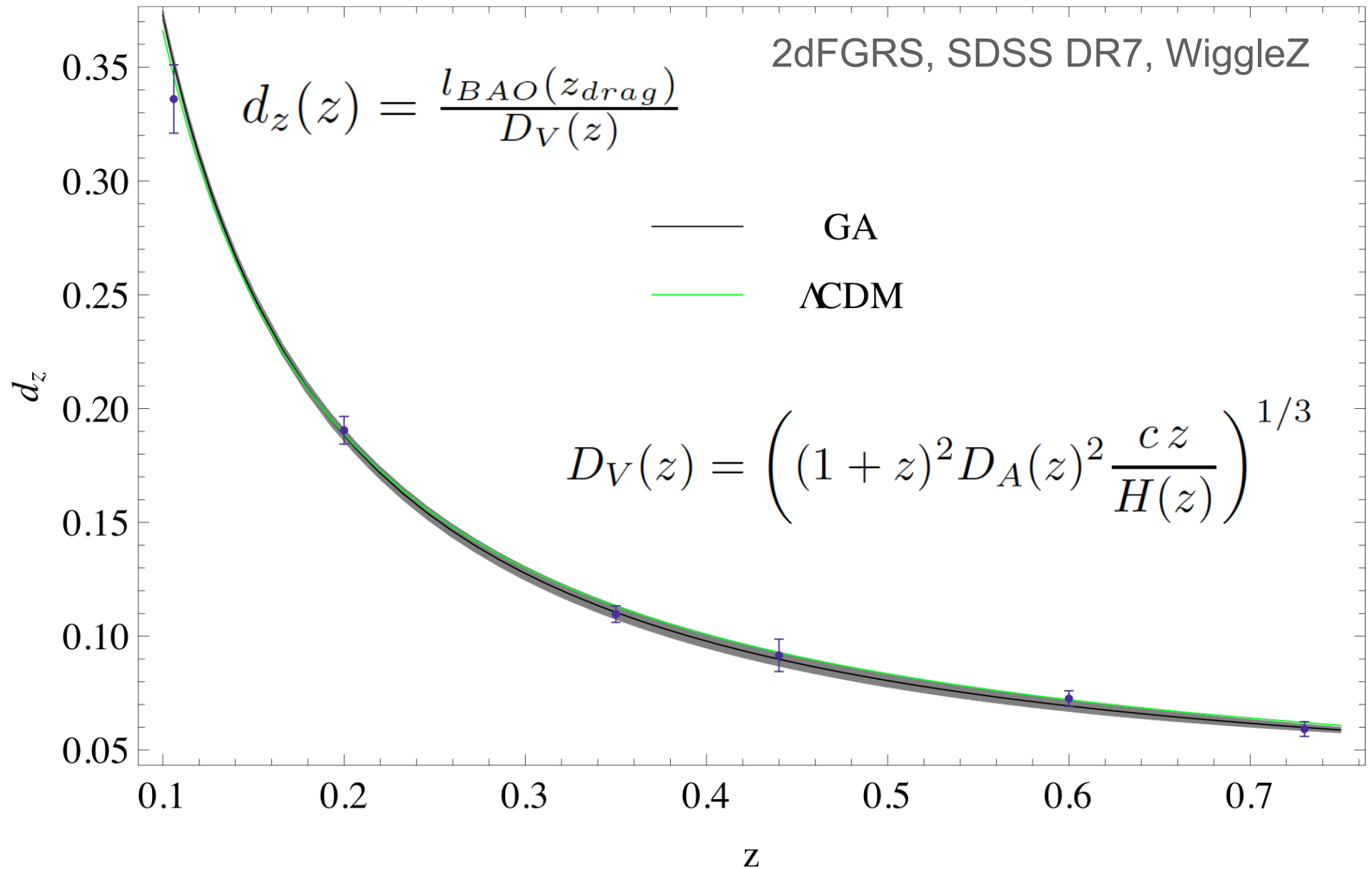
Union2.1, Suzuki et al. ApJ 746, 85 (2012)



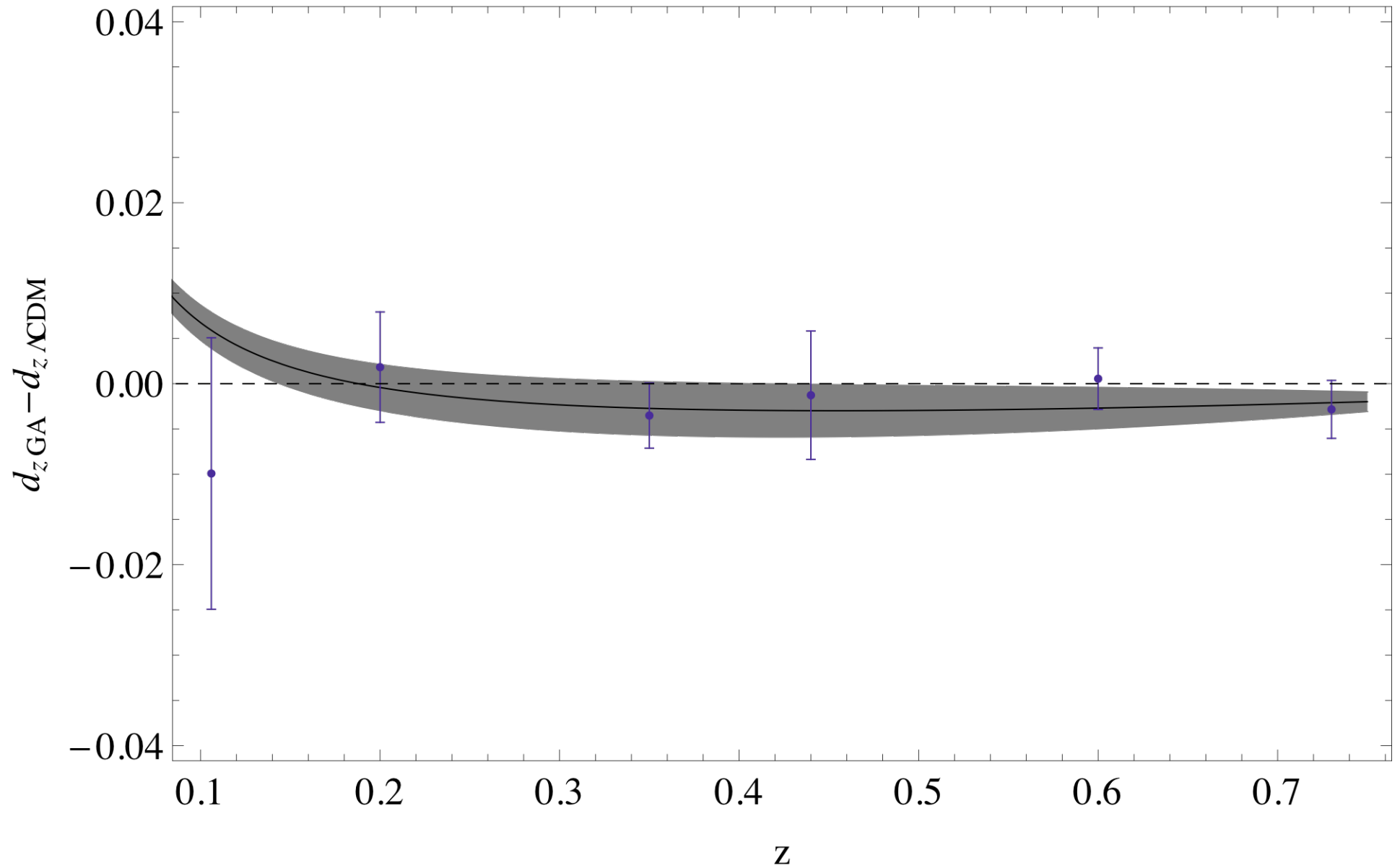
Genetic Algorithms & SNIa data



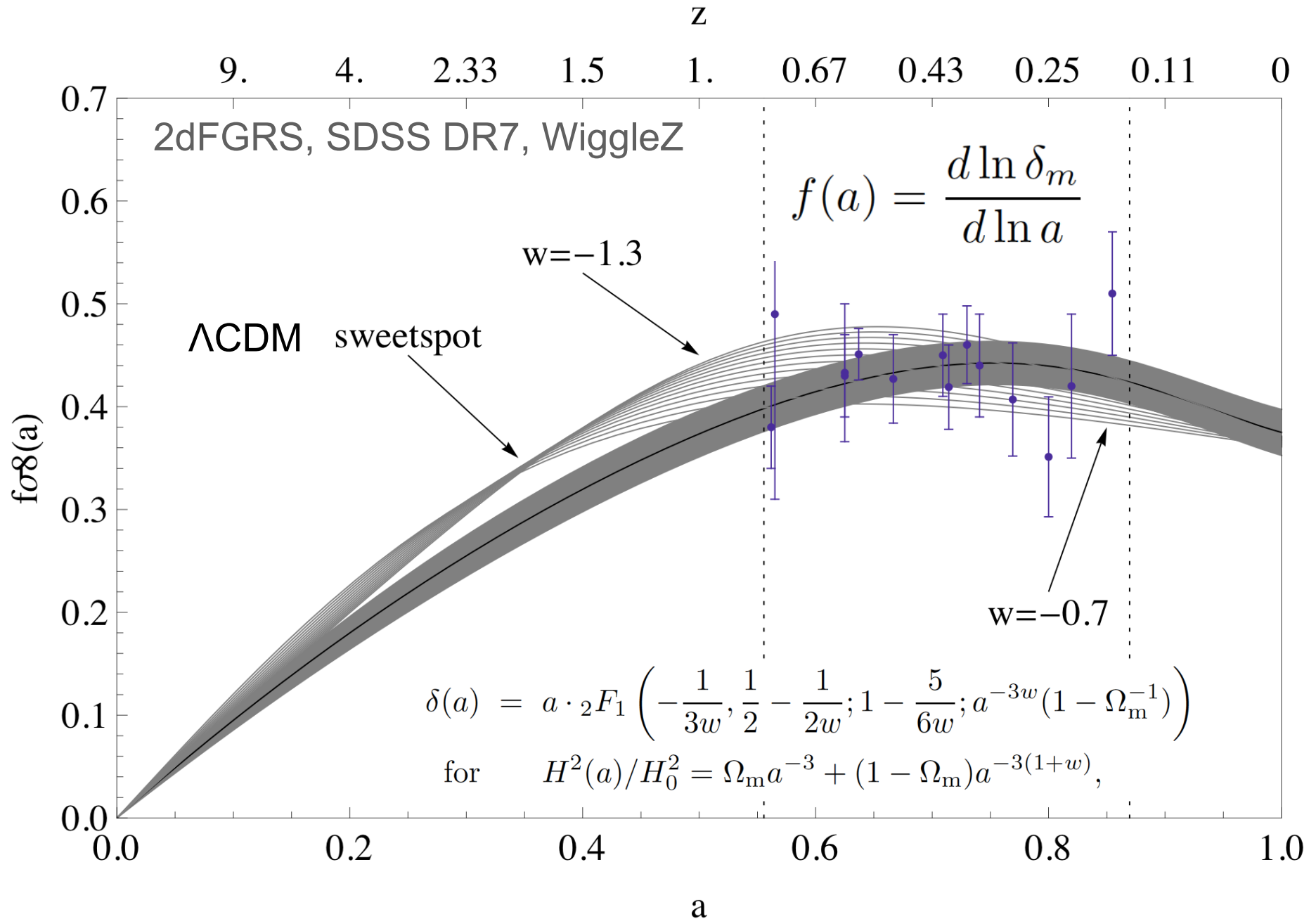
Genetic Algorithms & BAO data



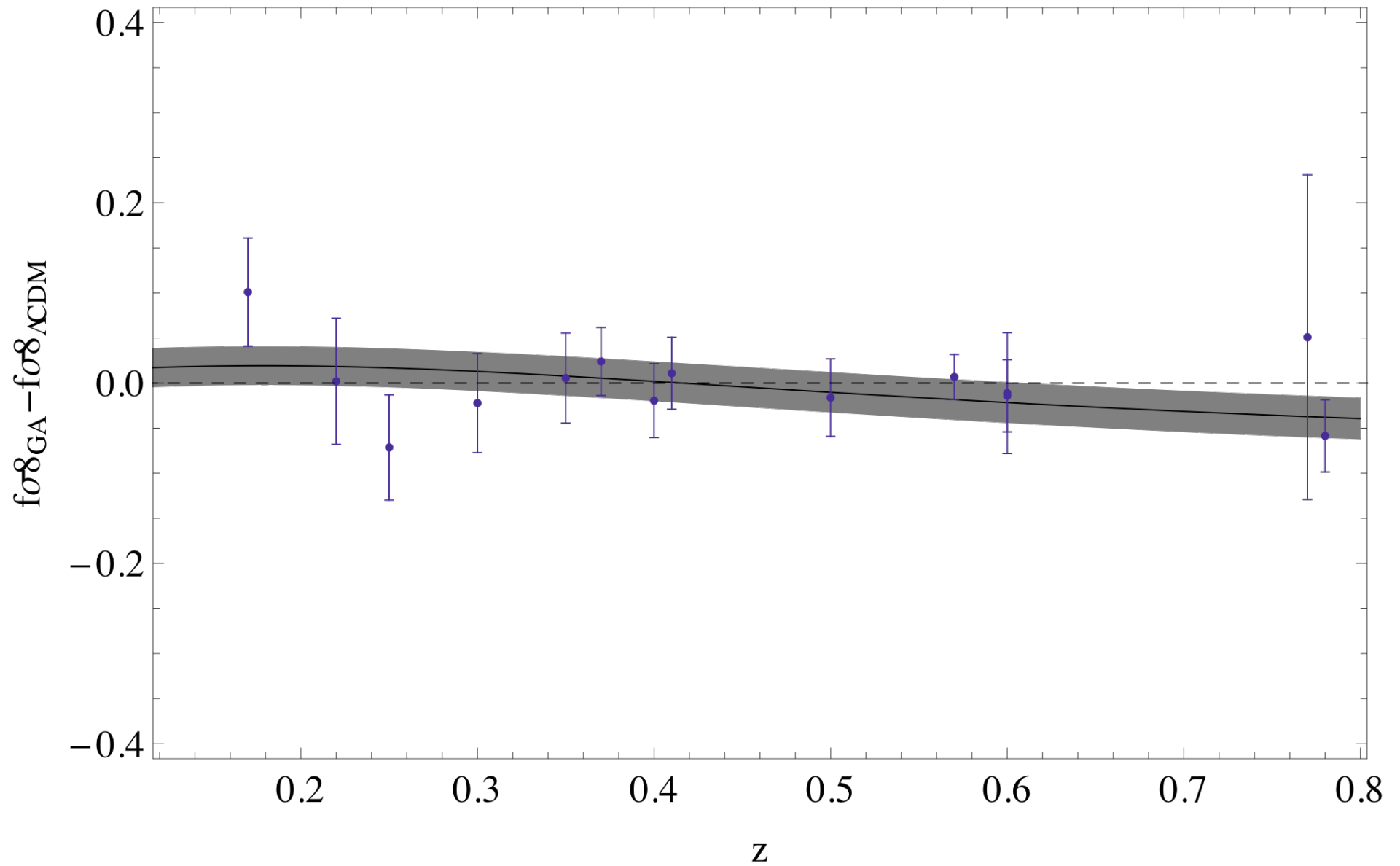
Genetic Algorithms & BAO data



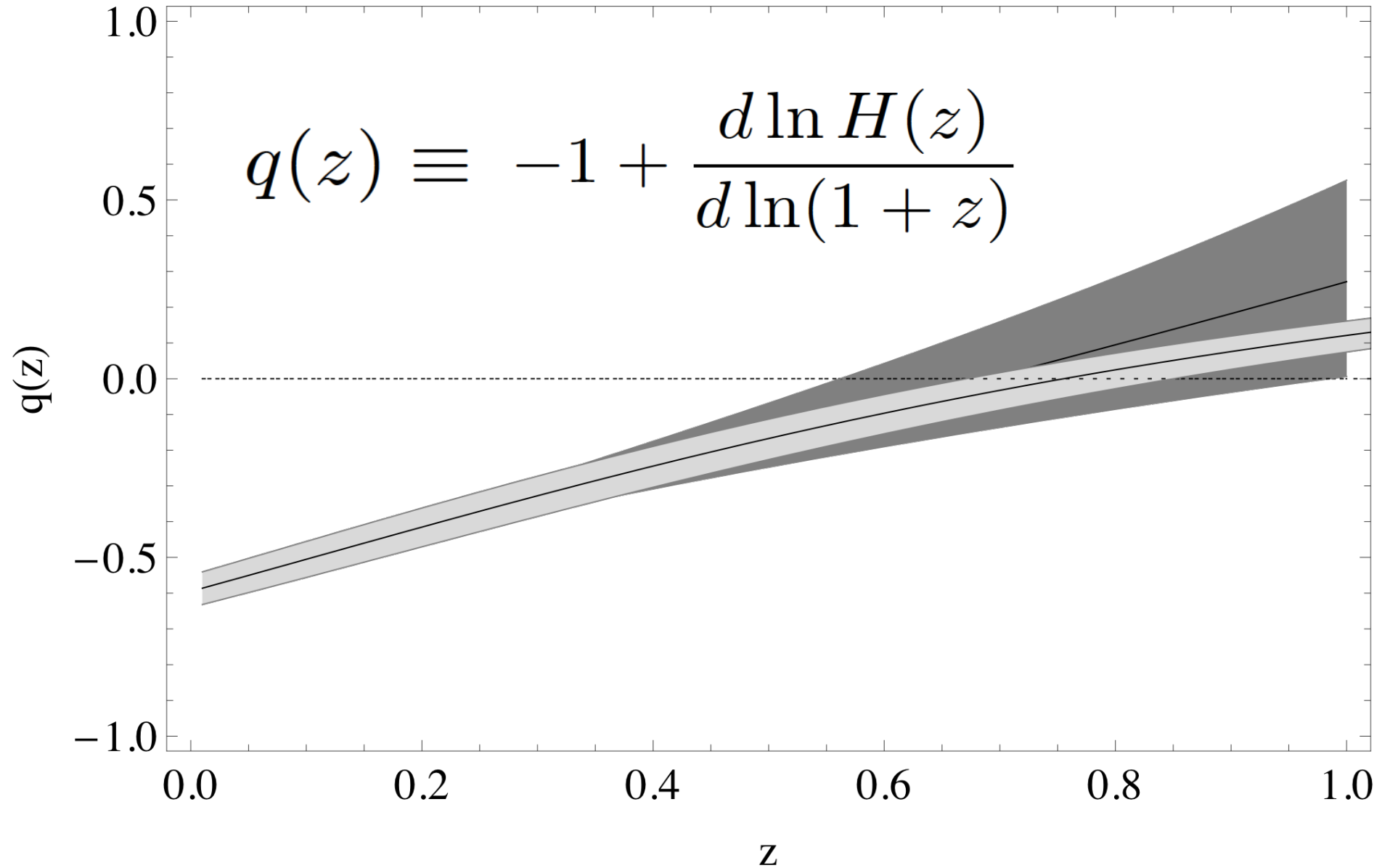
Genetic Algorithms & growth data



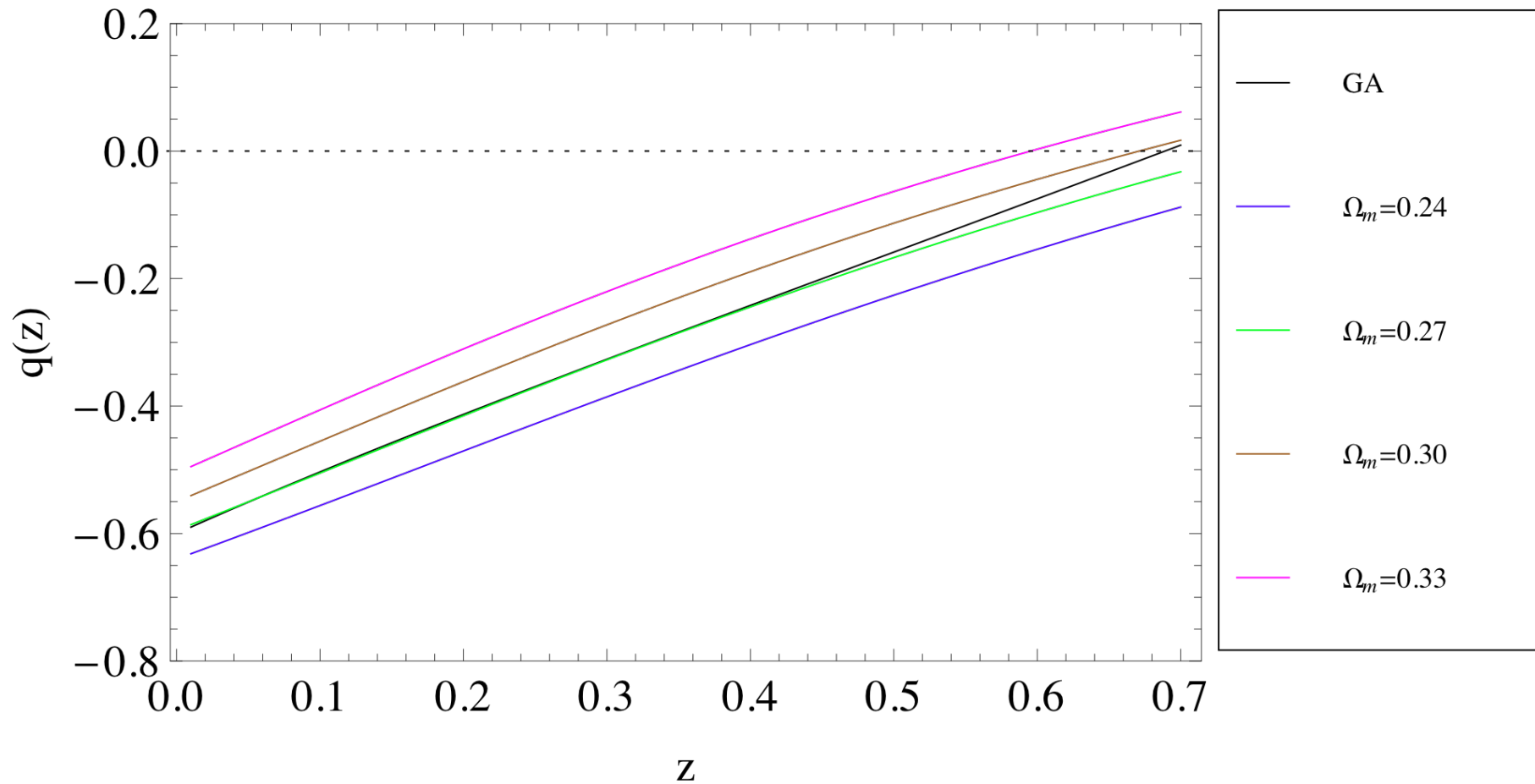
Genetic Algorithms & growth data



Genetic Algorithms & derived param.

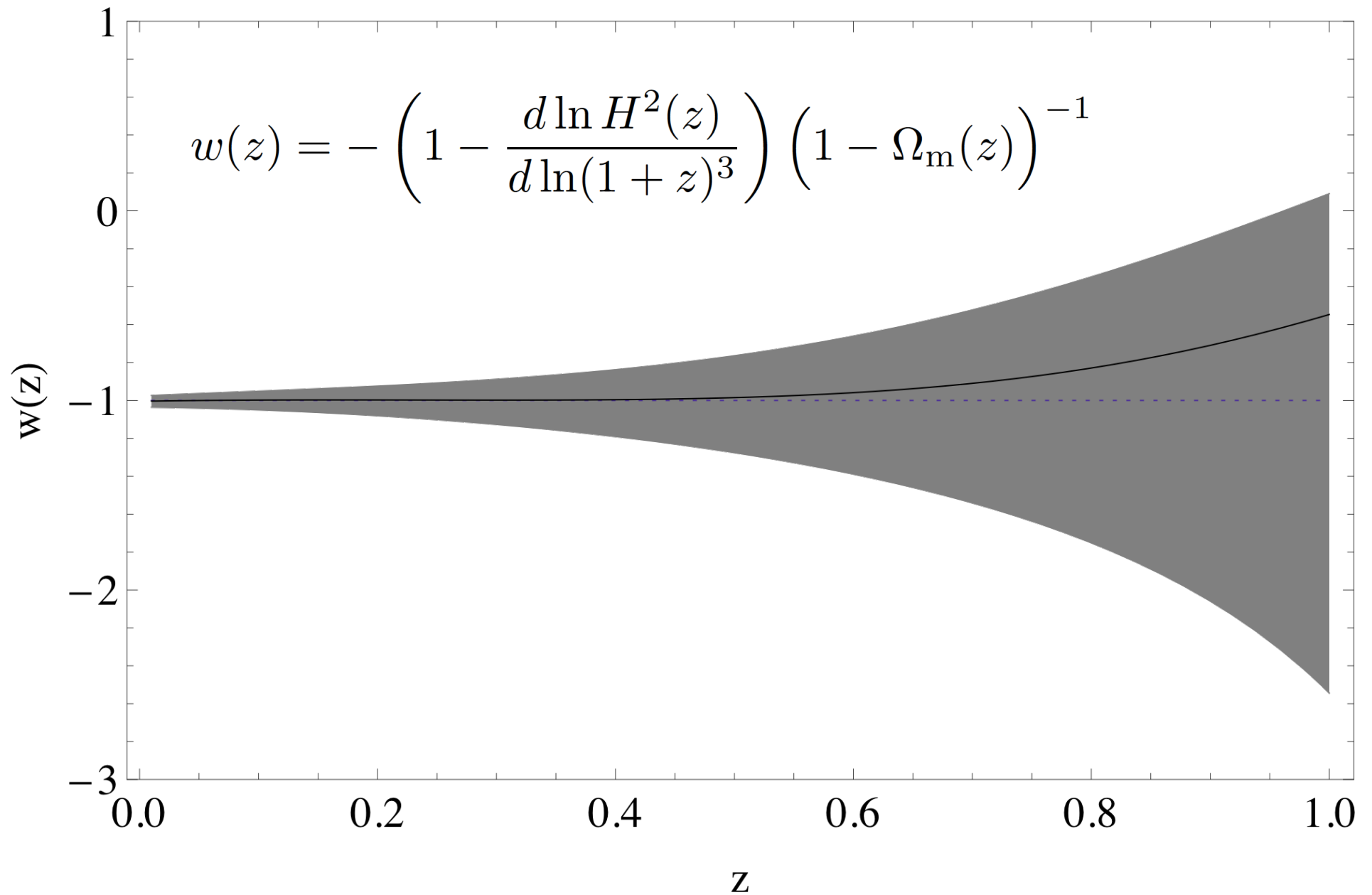


Genetic Algorithms & derived param.

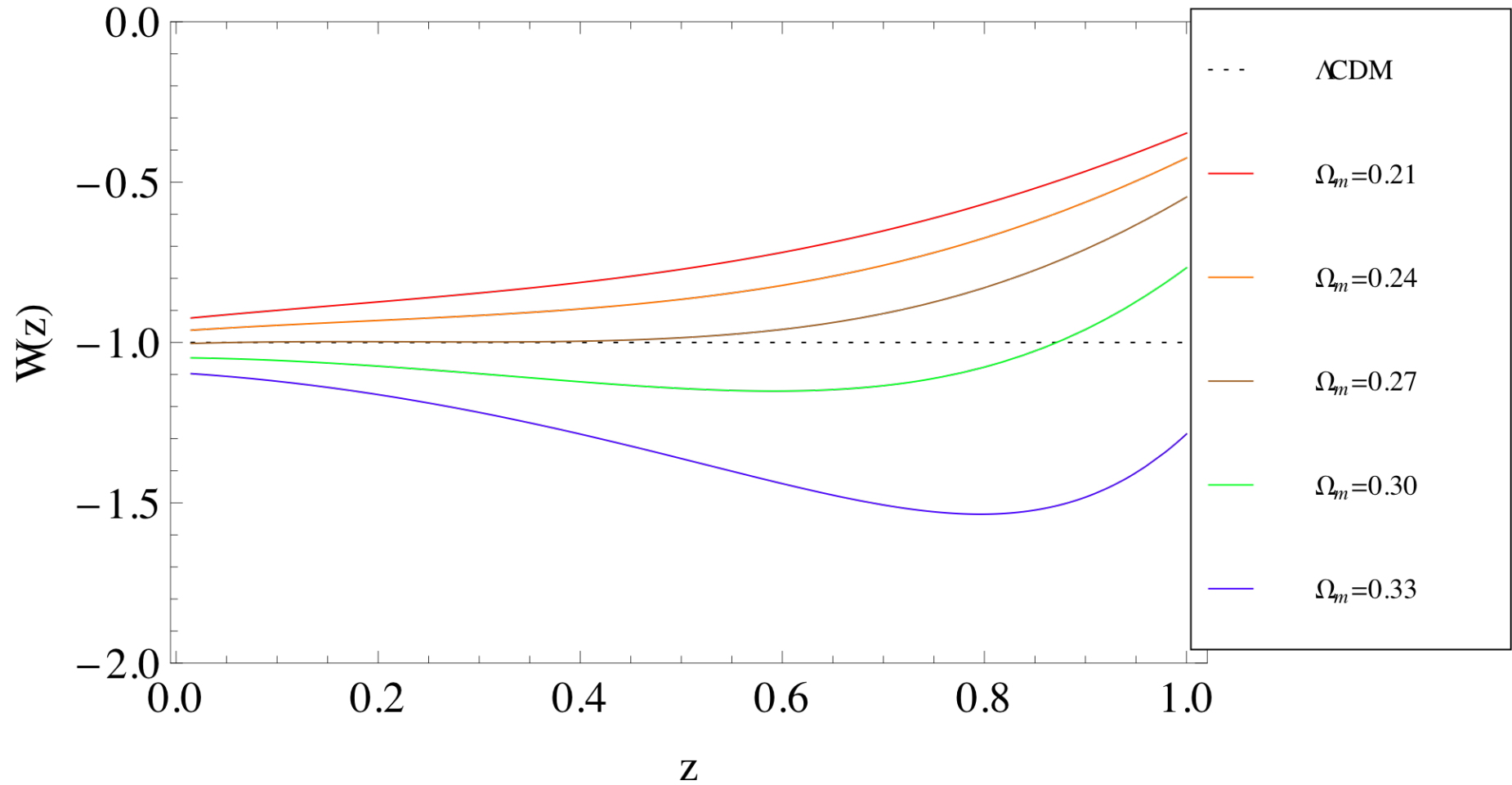


$$q(z) \equiv -1 + \frac{d \ln H(z)}{d \ln(1+z)}$$

Genetic Algorithms & derived param.

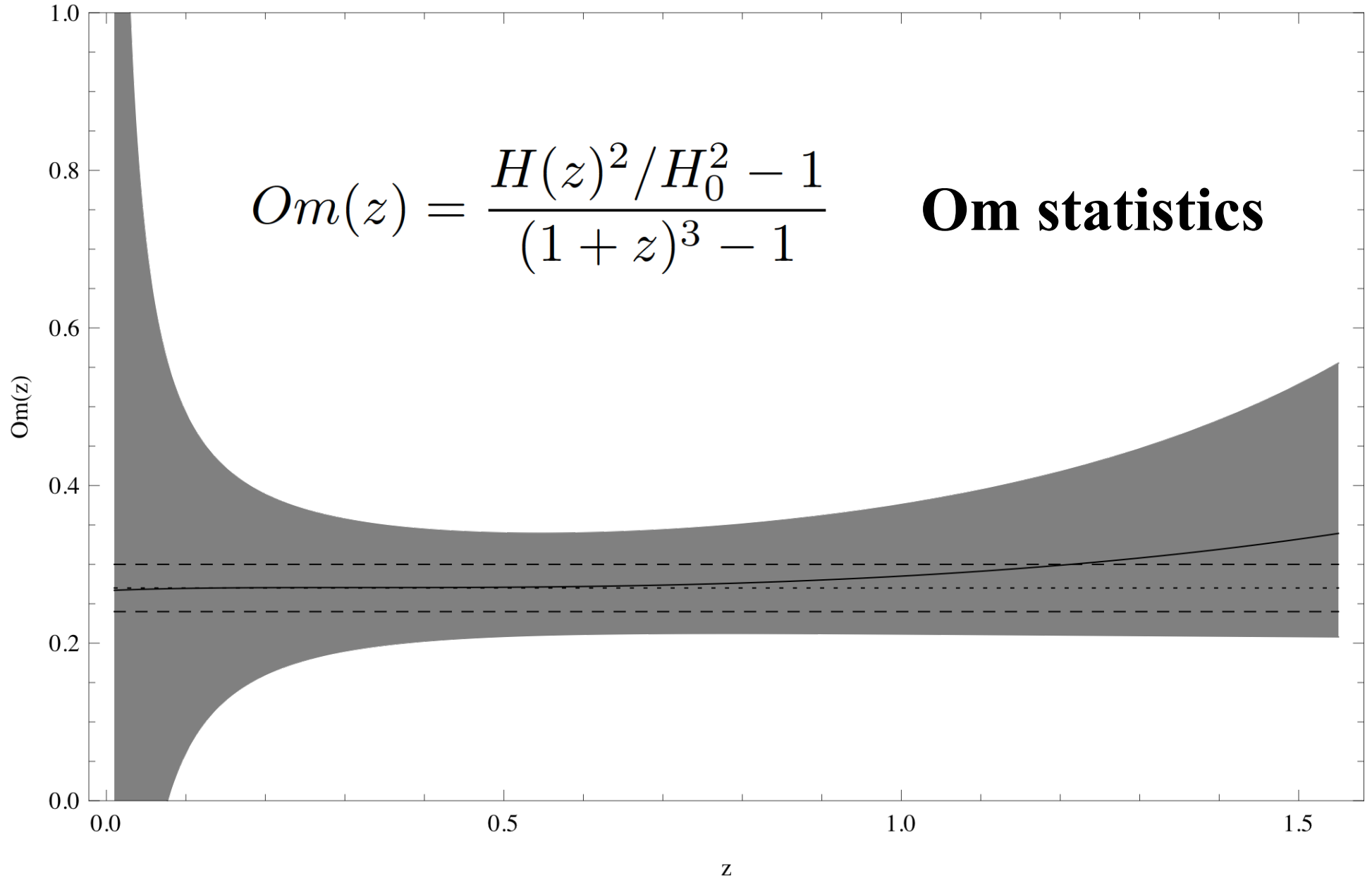


Genetic Algorithms & derived param.

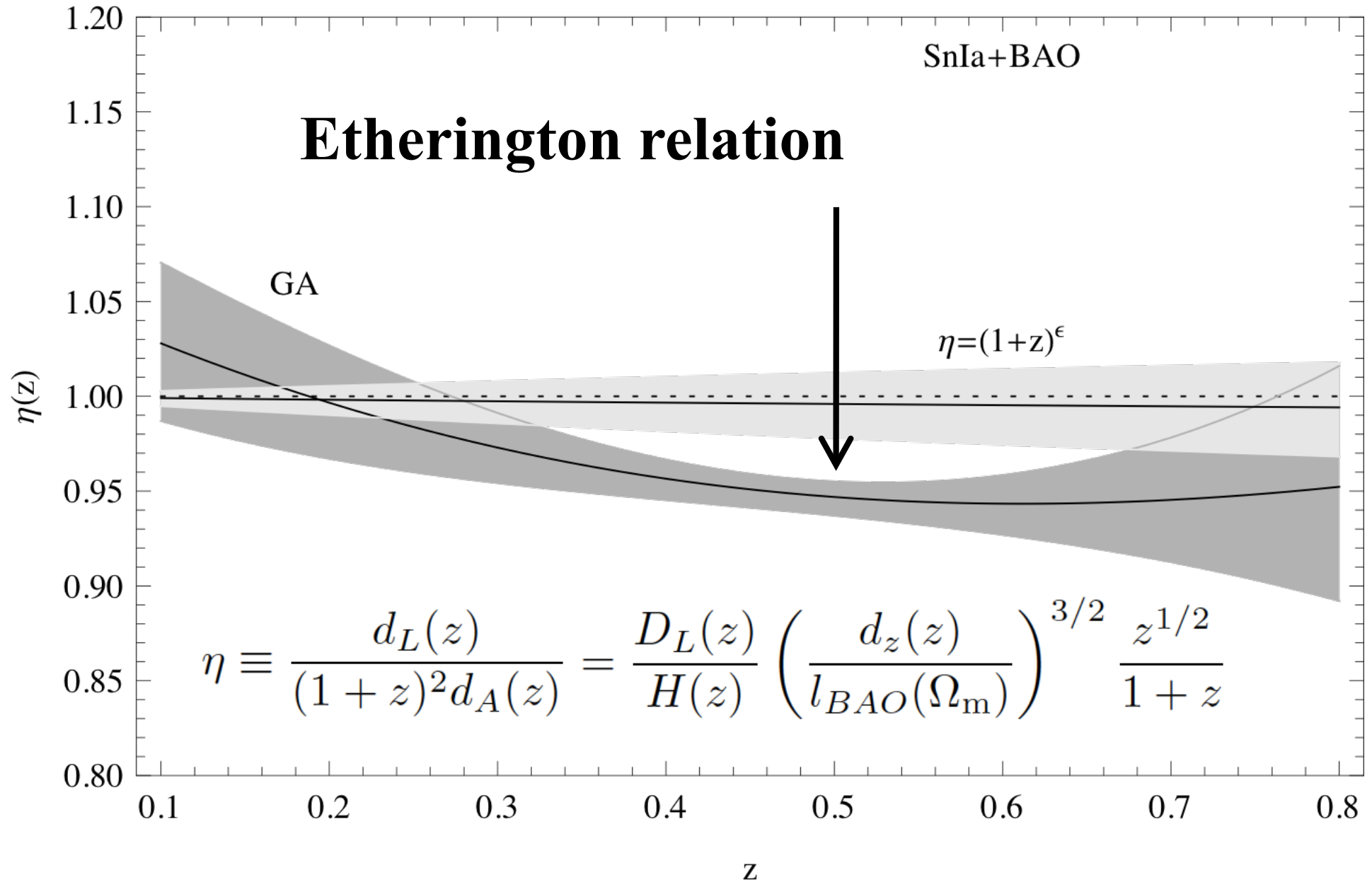


$$w(z) = - \left(1 - \frac{d \ln H^2(z)}{d \ln(1+z)^3} \right) \left(1 - \Omega_m(z) \right)^{-1}$$

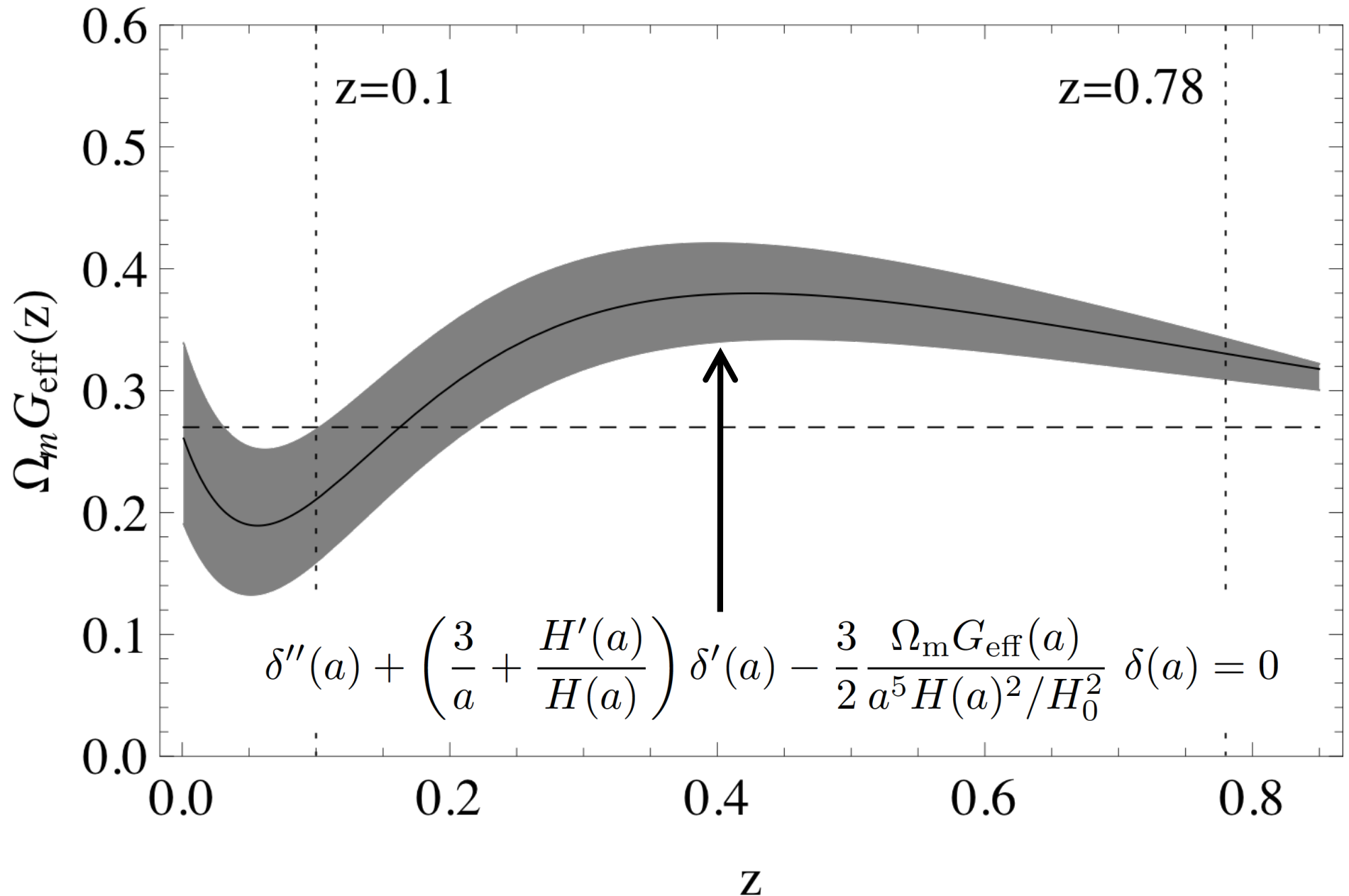
Genetic Algorithms & derived param.



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Conclusions

- **A decade since the discovery of acceleration, we still don't know what Dark Energy is!**
- **We need a model-independent and bias-free reconstruction of the observables that we deduce from data.**
- **We propose a new method: Genetic Algorithms**
- **Error estimation with Path Integral formalism**
- **Show how it works for SNIa, BAO & growth data**
- **Λ CDM works pretty well...**
- **Find slight deviations at 3σ on $\Omega_m G_{\text{eff}}$ @ $z=0.5$ and the Etherington relation**