# Microlensing: Theory, Practice, Results, Future Lecture 4

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Microlensing: Theory, Practice, Results, Future – p.1/30

Lecture 1: Microlensing History and Theory

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- Background
- Motivation/Goals
- Early results
- Evolution of a field
- Basic microlensing theory

- Lecture 1: Microlensing History and Theory
- Lecture 2: Beyond the Single lens

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- Lecture 2: Beyond the Single lens
  - Finite source star
  - Limb Darkening
  - Blending
  - Parallax
  - Xallarap

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- Lecture 3: Planetary Microlensing I

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- Lecture 2: Beyond the Single lens
- Lecture 3: Planetary Microlensing I
  - Binary lens microlensing
  - Extreme mass ratio microlensing
  - Theoretical tools of trade: caustics
  - Planetary microlensing regiemes
  - General rules

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- Lecture 4: Planetary Microlensing II

- Lecture 1: Microlensing History and Theory
- Lecture 2: Beyond the Single lens
- Lecture 3: Planetary Microlensing I
- Lecture 4: Planetary Microlensing II
  - Capabilities, detection limits
  - Detection
  - Modelling



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Microlensing is currently detecting planets in a previously unreachable region of the planetary mass-radius space. Microlensing is returning detections of planets with masses approaching that of Earth.

#### **Detection space**



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Distant planetary systems are being discovered.














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Ground-based microlensing is relatively insensitive to habitable planets, as the most-likely lens stars are cool M-dwarfs. However, a space telescope such as the proposed Microlensing Planet Finder (Bennett, 2006) will be much more likely to detect habitable planets via microlensing, including moons, if present.

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However:

In studies of nearby solar type stars, Abt & Levy (1976) found 58% of the G dwarf primary stars had one or more stellar companions( see also Abt, 1987). Duquennoy & Marcy (1991) find a similar fraction of 57%. The multiplicity amongst M dwarfs is slightly less, at 42% (Fischer & Marcy, 1992).

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Most common "goodness-of-fit" paramater is

$$\chi^{2} = \sum_{i=1}^{N} \frac{(y_{i}(t) - \hat{y}_{i}(t))^{2}}{\sigma_{i}^{2}}$$

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Detection limit maps for an Earth mass planet orbiting a  $0.3M_{\odot}$  lens star. The axes are in units of  $R_E$ . The I band magnitude at maximum is 15, and the maximum amplification is 100. The  $\chi^2$  map on the left was generated using observation light curves comprised of 301 points over the interval  $\left[-\frac{1}{2}t_{\rm FWHM}, \frac{1}{2}t_{\rm FWHM}\right]$ . The right map was created using light curves comprised of 601 points over the interval  $\left[-t_{\rm FWHM}, t_{\rm FWHM}\right]$ .

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- Point-by-point significance e.g 3 consecutive points deviating by  $\geq 3\sigma$
- Coherency deviating points follow a clear trend
- Confirmable are deviating points supported by more observations

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But before we can apply detection criteria, we need to be able to fit microlensing lightcurve models!

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The position of a planet is subject to a further degeneracy: models with projected planet distance  $a_p$  are degenerate with those with distance  $1/a_p$ .

Point source single lens events are relatively easy to model. Standard non-linear fitting algorithms can be used to find  $u_{\min}$ ,  $t_E$  and  $t_0$ . The blend flux parameters  $F_u$ ,  $F_l$  can similarly be found, either as additional parameters in the non-linear fitting routine, or through linear least-squares.

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$$u(t) = \left[ u_{\min}^{2} + \left( \frac{(t - t_{0})}{t_{E}} \right)^{2} \right]^{\frac{1}{2}}$$
$$\mu = \frac{u^{2} + 2}{u\sqrt{u^{2} + 4}}$$

 $F = F_1 \cdot A(u(t)) + F_{\text{Microlensing: Theory, Practice, Results, Future - p.16/3}$ 

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Inverse ray shooting is a numerical technique which, although slow, is robust:

• (Rattenbury, 2002)

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The fitting algorithm used also needs to be well understood.

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The first coarse search could be a grid-wise search for example, or a random (uniform) sampling over the parameter space.

May need several restarts, but should provide some good starting locations.



Having determined where possible optimal parameter values might lie in the parameter space  $\Omega = \Omega\{u_{\min}, t_0, t_E, r_s, x_{p_i}, y_{p_i}, q_i\}$ , the next step is to apply more rigorous  $\chi^2$  minimisation procedures.

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 $\chi^2$  minima.

The simplex method always moving toward a lower  $\chi^2$  value. If the  $\chi^2$  manifold over the N-dimensional parameter space is not smooth, methods such as the downhill simplex can get trapped in non-optimal parameter space states.

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It is for this reason that it is suggested that the simplex method is restarted often, around the point in  $\Omega$  considered to be the set of parameter values that minimises  $\chi^2$ . If the procedure recovers the same point after starting from a number of different points, the set of optimal parameters is considered secure.

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If the  $\chi^2$  manifold over  $\Omega$  is sufficiently complicated, there is a greater risk that the results returned from the simplex methods are not global minima.

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This feature of the MHMCMC algorithm allows the method to occasionally disregard a (possibly better) candidate state. A chance therefore exists for the algorithm to find a more favourable region on the  $\chi^2$  manifold.



Assume that the current set of parameters is  $\Omega$ .

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Let  $\{Z_k\}_{k=1}^M$  define a Markov chain, length M. Each element of  $Z_k$  is a set of parameters:  $Z_k = \Omega_k$ .

The next element in the chain,  $Z_{k+1}$  is determined as follows:

1. Choose a candidate set of parameters,  $\Omega$ ', by varying one or more of the parameters:

$$u_{\min} ' = u_{\min} + \mathcal{R}_{u_{\min}} \qquad (0)$$

$$t'_{0} = t_{0} + \mathcal{R}_{t_{0}}$$

$$t'_{E} = t_{E} + \mathcal{R}_{t_{E}}$$

$$\vdots = \vdots$$

where  $\mathcal{R}$  is an appropriately scaled random number.

2. The candidate state,  $\Omega$ ', is accepted with probability:

 $\alpha(\mathbf{\Omega}'|\mathbf{\Omega}) = \min\left\{1, \mathcal{F}(f'(t), f(t))\right\}$ 

where f(t) is the model function calculated using the current parameter set,  $\Omega$ . f'(t) is the model function, calculated using the candidate parameter set,  $\Omega'$ .  $\mathcal{F}$  is a function based on the difference between the current and candidate models. If the candidate state  $\Omega'$  is not accepted, set  $Z_{k+1} = \Omega_k$ .

 $\alpha(\Omega'|\Omega)$  is the acceptance ratio. The function  $\mathcal{F}$  essentially compares the  $\chi^2$  values arising from fitting the light curve generated using the candidate parameters  $\Omega$ ', to that of the current set.

$$\mathcal{F} = \exp\left[\frac{\chi^2 - \chi'^2}{2}\right]$$

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The above two procedures are iterated M times. Each parameter is allowed to vary, and the algorithm settles at a point in the parameter space which minimises  $\chi^2$ .



We can also impose an annealing schedule, T(n):

$$\mathcal{F} = \exp\left[\frac{\chi^2 - \chi'^2}{2T}\right]$$

If  $T \rightarrow 0$ , the distribution will converge on the optimal modal space in the state space. Given a distribution in equilibrium, the "temperature", T, can slowly be lowered and allow the distribution to find its new equilibrium. As  $T \rightarrow 0$  the equilibrium will settle on the mode state(s).

	$\Delta \chi^2 > 0$		$\Delta \chi^2 < 0$	
T	lpha(z' z)	Accept $z'$	lpha(z' z)	Accept $z'$
> 1	1	always	$\leq 1$	often
1.0	1	always	$0 < \alpha < 1$	sometimes
0 < T < 1	1	always	$0 < \alpha \ll 1$	seldom
0.0	1	always	0	never
$\Delta \chi^2 = \chi^2 - \chi'^2$ , where $\chi'^2$ corresponds to the				
candidate state, and $\chi^2$ to the current state.				

- MHMCMC is excellent for fitting microlensing lightcurves
- MHMCMC fitting can be optimised in many ways.
- MHMCMC can be parallized easily.
- There is a lot of information in the equilibrium state of each parameter.

Genetic algorithms such as Particle Swarm Optimisation, Artificial Neural Networks and Self Organising Maps are other possible categorization methods.